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JC526 U.S. PTO
09/198779
11/24/98

November 24, 1998

Box Patent Application

Assistant Commissioner for Patents
Washington, D.C. 20231

Re: U.S. Non-Provisional Patent Application
Appl. No.: To Be Assigned
Filed: November 24, 1998
For: **NUCLEIC ACID MOLECULES AND OTHER MOLECULES
ASSOCIATED WITH THE METHIONINE SYNTHESIS AND
DEGRADATION PATHWAYS**
Inventor(s): Stefan A. Bledig *et al.*
Our Ref: 04983.0002US01/38-21 (15077)B

Sir:

The following documents are being submitted under 37 C.F.R. § 1.53(b)(2) herewith for appropriate action by the U.S. Patent and Trademark Office:

1. Patent Application Utility Application;
2. Utility Patent Application Transmittal (PTO/SB/05);
3. Form PTO-1082 (in duplicate);
4. U.S. Utility Patent Application entitled

**Nucleic Acid Molecules And Other Molecules Associated With The
Methionine Synthesis And Degradation Pathways**

and naming as inventor(s):

**Stefan A. Bledig, Joseph R. Byrum, Gregory J. Hinkle and JingDong
Liu**

the application consisting of:

- (i) 288 pages of description prior to the claims;
 - (ii) 5 pages of claims (11 claims);
 - (iii) a one (1) page abstract;
 - (iv) sequence listing of 1127 pages;
- 5. Sequence Listing Disk;
 - 6. Statement Regarding Sequence Submissions;
 - 7. Information Disclosure Statement;
 - 8. Form PTO-1449 (9 pages) with 30 accompanying documents; and
 - 9. Two (2) return postcards.

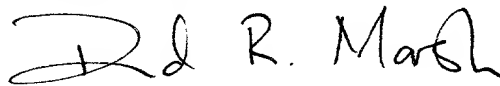
It is respectfully requested that, of the two attached postcards, one be stamped with the filing date of these documents and returned to our courier, and the other, prepaid postcard, be stamped with the filing date and returned as soon as possible.

The present application claims priority under 35 U.S.C § 119(e) to provisional applications No. 60/067000 filed November 24, 1997, No. 60/066,873 filed November 25, 1997, No. 60/069472 filed December 09, 1997, No. 60/074,201 filed February 10, 1998, No. 60/074282 filed February 10, 1998, No. 60/074280 filed February 10, 1998, No. 60/074281 filed February 10, 1998, No. 60/074566 filed February 12, 1998, No. 60/074567 filed February 12, 1998, No. 60/074565 filed February 12, 1998, No. 60/075462 filed February 19, 1998, No. 60/074789 filed February 19, 1998, No. 60/075459 filed February 19, 1998, No. 60/075461 filed February 19, 1998, No. 60/075464 filed February 19, 1998, No. 60/075460 filed February 19, 1998, No. 60/075463 filed February 19, 1998, No. 60/077231 filed March 09, 1998, No. 60/077229 filed March 09, 1998, No. 60/077230 filed March 09, 1998, No. 60/078031 filed March 16, 1998, No. 60/078368 filed March 18, 1998, No. 60/080844 filed April 07, 1998, No. 60/083067 filed April 27, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15348)A filed April 29, 1998, No. 60/083387 filed April 29, 1998, No. 60/083388 filed April 29, 1998, No. 60/083389 filed April 29, 1998, "Nucleic Acid Molecules and Other Molecules Associated with the Ethylene Biosynthetic Pathway" docket No. 04983.0018/38-21(15097)A filed May 08, 1998, No. 60/085,245 filed May 13, 1998, No.

60/085224 filed May 13, 1998, No. 60/085223 filed May 13, 1998, No. 60/085222 filed May 13, 1998, No. 60/086186 filed May 21, 1998, No. 60/086,339 filed May 21, 1998, No. 60/086187 filed May 21, 1998, No. 60/086185 filed May 21, 1998, No. 60/086184 filed May 21, 1998, No. 60/086183 filed May 21, 1998, No. 60/086188 filed May 21, 1998, No. 60/089,524 filed June 16, 1998, No. 60/089,810 filed June 18, 1998, No. 60/089,814 filed June 18, 1998, "Nucleic acid molecules and other molecules associated with the Plant Sugar and Nitrogen Transporters Pathway" docket No. 04983.0043/38-21(15412)A filed June 30, 1998, No. 60/092,036 filed July 08, 1998, No. 60/099667 filed September 09, 1998, No. 60/099668 filed September 09, 1998, No. 60/099670 filed September 09, 1998, No. 60/099697 filed September 09, 1998, No. 60/100674 filed September 16, 1998, No. 60/100673 filed September 16, 1998, No. 60/100672 filed September 16, 1998, No. 60/101132 filed September 21, 1998, No. 60/101130 filed September 21, 1998, "Nucleic acid molecules and other molecules associated with Plants" docket No. 38-21(15459)A filed September 21, 1998, No. 60/101344 filed September 22, 1998, No. 60/101347 filed September 22, 1998, No. 60/101343 filed September 22, 1998, No. 60/104,126 filed October 13, 1998, No. 60/104,128 filed October 13, 1998, No. 60/104,127 filed October 13, 1998, No. 60/104,124 filed October 13, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15445)A filed November 18, 1998 and "Nucleic Acid Molecules and other Molecules associated with Plants" docket No. 38-21(15592) filed November 18, 1998 hereby incorporated by reference herein in their entirety.

This application is being filed under 37 C.F.R. § 1.53(b) without Declaration and without filing fee.

Respectfully submitted,



David R. Marsh (Reg. No. 41,408)

Kevin W. McCabe (Reg. No. 41,182)

Enclosures

Please type a plus sign (+) inside this box → ☐

UTILITY PATENT APPLICATION TRANSMITTAL <small>(Only for new nonprovisional applications under 37 CFR 1.53(h))</small>		Attorney Docket No. 04983.0002US01/38-21(15077)B	
		First Named Inventor or Application Identifier Stefan A. Bledig	
		Title Nucleic Acid Molecules and Other Molecules Associated with the Methionine Synthesis and Degradation Pathways	
		Express Mail Label No.	

APPLICATION ELEMENTS <i>See MPEP chapter 600 concerning utility patent application contents</i>	ADDRESS TO: Assistant Commissioner for Patents Box Patent Application Washington, DC 20231
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1. ☒ ⁴ Fee Transmittal Form (Form PTO-1082)
(Submit an original and a duplicate for fee processing)

2. ☒ Specification [Total Pages 294]
(preferred arrangement set forth below)
- Descriptive title of the Invention
- Cross References to Related Applications
- Statement Regarding Fed sponsored R&D
- Reference to Microfiche Appendix
- Background of the Invention
- Brief Summary of the Invention
- Brief Description of the Drawings (if filed)
- Detailed Description
- Claims
- Abstract of the Disclosure

3. ☐ Drawing(s) (35 USC 113) [Total Sheets]

4. ☐ Oath or Declaration [Total Pages]

☐ Newly executed (original or copy)
☐ Copy from a prior application (37 CFR 1.63(d))
(for continuation/divisional with Box 17 completed)
[Note Box 5 below]

☐ **DELETION OF INVENTOR(S)**
Signed statement attached deleting inventor(s) named in the prior application, see 37 CFR 1.63(d)(2) and 1.33(b)

5. ☐ Incorporation By Reference *(useable if Box 4b is checked)*
The entire disclosure of the prior application, from which a copy of the oath or declaration is supplied under Box 4b, is considered as being part of the disclosure of the accompanying application and is hereby incorporated by reference therein.

6. ☐ Microfiche Computer Program *(Appendix)*

7. Nucleotide and/or Amino Acid Sequence Submission *(if applicable, all necessary)*

☒ Computer Readable Copy
☒ Paper Copy (identical to computer copy)
☒ Statement verifying identity of above copies

ACCOMPANYING APPLICATION PARTS

8. ☐ Assignment Papers (cover sheet & document(s))

9. ☐ 37 CFR 3.73(b) Statement ☐ Power of Attorney
(when there is an assignee)

10. ☐ English Translation Document *(if applicable)*

11. ☒ Information Disclosure Statement (IDS)/PTO-1449 ☒ Copies of IDS Citations

12. ☐ Preliminary Amendment

13. ☒ Return Receipt Postcard (MPEP 503) (Two)
(should be specifically itemized)

14. ☐ *Small Entity Statement(s) ☐ Statement filed in prior application, Status still proper and desired

15. ☐ Certified Copy of Priority Document(s)
(if foreign priority is claimed)

16. ☐ Other:

*NOTE FOR ITEMS 1 & 14 IN ORDER TO BE ENTITLED TO PAY SMALL ENTITY FEES, A SMALL ENTITY STATEMENT IS REQUIRED (37 C.F.R. § 1.27). EXCEPT IF ONE FILED IN A PRIOR APPLICATION IS RELIED UPON (37 C.F.R. § 1.28)

17. **If a CONTINUING APPLICATION.** check appropriate box and supply the requisite information:
☐ Continuation ☐ Divisional ☐ Continuation-in-part (CIP) of prior application No: /
Prior Application Information: Examiner: Group/Art Unit:

18. CORRESPONDENCE ADDRESS

☐ Customer Number or Bar Code Label
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or ☒ Correspondence address below

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Attorney Docket No. 04983.0002US01/38-21 (15077)B

ASSISTANT COMMISSIONER FOR PATENTS
 Washington, D.C. 20231

Sir:

Transmitted herewith for filing is the patent application of
 Inventors: Stefan A. Bledig *et al.*

For: **Nucleic Acid Molecules and Other Molecules Associated with the Methionine Synthesis and Degradation Pathways**

The filing fee has been calculated as shown below:

	(Col. 1)	(Col. 2)
FOR	NO. FILED	NO EXTRA
BASIC FEE		
TOTAL CLAIMS	-20 =	*
INDEP. CLAIMS	-3 =	*
MULTIPLE DEPENDENT CLAIM PRESENTED		

SMALL ENTITY	
RATE	FEE
	\$ 380.00
x 9 =	
x 39 =	
+ 130 =	
TOTAL	

OTHER THAN A SMALL ENTITY	
RATE	FEE
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x 18 =	
x 78 =	
+ 260 =	
TOTAL	

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The U.S. Patent and Trademark Office is hereby authorized to charge payment of the following fees associated with this communication or credit any overpayment to Deposit Account No. 08-3038. A duplicate copy of this sheet is attached.

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The U.S. Patent and Trademark Office is hereby authorized to charge payment of the following fees during the pendency of this application or credit any overpayment to Deposit Account No. 08-3038. A duplicate copy of this sheet is enclosed.

- Any patent application processing fees under 37 CFR 1.17
 — The issue fee set in 37 CFR 1.18 at or before mailing of the Notice of Allowance, pursuant to 37 CFR 1.311(b).
 — Any filing fees under 37 CFR 1.16 for presentation of extra claims.

Date November 24, 1998



David R. Marsh (Reg. No. 41,408)
 Kevin W. McCabe (Reg. No. 41,182)

NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH

THE METHIONINE SYNTHESIS AND DEGRADATION PATHWAYS

CROSS REFERENCE TO RELATED APPLICATIONS

This application claims the benefit of applications No. 60/067000 filed November 24, 1997, No. 60/066,873 filed November 25, 1997, No. 60/069472 filed December 09, 1997, No. 60/074,201 filed February 10, 1998, No. 60/074282 filed February 10, 1998, No. 60/074280 filed February 10, 1998, No. 60/074281 filed February 10, 1998, No. 60/074566 filed February 12, 1998, No. 60/074567 filed February 12, 1998, No. 60/074565 filed February 12, 1998, No. 60/075462 filed February 19, 1998, No. 60/074789 filed February 19, 1998, No. 60/075459 filed February 19, 1998, No. 60/075461 filed February 19, 1998, No. 60/075464 filed February 19, 1998, No. 60/075460 filed February 19, 1998, No. 60/075463 filed February 19, 1998, No. 60/077231 filed March 09, 1998, No. 60/077229 filed March 09, 1998, No. 60/077230 filed March 09, 1998, No. 60/078031 filed March 16, 1998, No. 60/078368 filed March 18, 1998, No. 60/080844 filed April 07, 1998, No. 60/083067 filed April 27, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15348)A filed April 29, 1998, No. 60/083387 filed April 29, 1998, No. 60/083388 filed April 29, 1998, No. 60/083389 filed April 29, 1998, "Nucleic Acid Molecules and Other Molecules Associated with the Ethylene Biosynthetic Pathway" docket No. 04983.0018/38-21(15097)A filed May 08, 1998, No. 60/085,245 filed May 13, 1998, No. 60/085224 filed May 13, 1998, No. 60/085223 filed May 13, 1998, No. 60/085222 filed May 13, 1998, No. 60/086186 filed May 21, 1998, No. 60/086,339 filed May 21, 1998, No. 60/086187 filed May 21, 1998, No. 60/086185 filed May 21, 1998, No. 60/086184 filed May 21, 1998, No. 60/086183 filed May 21, 1998, No. 60/086188 filed May 21, 1998, No. 60/089,524 filed June 16, 1998, No. 60/089,810 filed June 18, 1998, No.

60/089,814 filed June 18, 1998, "Nucleic acid molecules and other molecules associated with the Plant Sugar and Nitrogen Transporters Pathway" docket No. 04983.0043/38-21(15412)A filed June 30, 1998, No. 60/092,036 filed July 08, 1998, No. 60/099667 filed September 09, 1998, No. 60/099668 filed September 09, 1998, No. 60/099670 filed September 09, 1998, No. 60/099697 filed September 09, 1998, No. 60/100674 filed September 16, 1998, No. 60/100673 filed September 16, 1998, No. 60/100672 filed September 16, 1998, No. 60/101132 filed September 21, 1998, No. 60/101130 filed September 21, 1998, "Nucleic acid molecules and other molecules associated with Plants" docket No. 38-21(15459)A filed September 21, 1998, No. 60/101344 filed September 22, 1998, No. 60/101347 filed September 22, 1998, No. 60/101343 filed September 22, 1998, No. 60/104,126 filed October 13, 1998, No. 60/104,128 filed October 13, 1998, No. 60/104,127 filed October 13, 1998, No. 60/104,124 filed October 13, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15445)A filed November 18, 1998 and "Nucleic Acid Molecules and other Molecules associated with Plants" docket No. 38-21(15592) filed November 18, 1998 hereby incorporated by reference herein in their entirety.

FIELD OF THE INVENTION

The present invention is in the field of plant biochemistry. More specifically the invention relates to nucleic acid sequences from plant cells, in particular, DNA sequences from maize and soybean plants associated with the methionine pathway. The invention encompasses nucleic acid molecules that encode proteins and fragments of proteins. In addition, the invention also encompasses proteins and fragments of proteins so encoded and antibodies capable of binding these proteins or fragments. The invention also relates to methods of using the nucleic acid molecules, proteins and fragments of proteins and antibodies, for example for genome

mapping, gene identification and analysis, plant breeding, preparation of constructs for use in plant gene expression and transgenic plants.

BACKGROUND OF THE INVENTION

I. METHIONINE SYNTHESIS PATHWAY

The amino acid, L-methionine, is synthesized in higher plants via a pathway that starts with L-aspartate. This pathway has been studied (Azevedo *et al.*, *Phytochemistry* 46:395-419 (1997), the entirety of which is herein incorporated by reference). L-methionine is one of four so-called aspartate-derived amino acids (along with L-lysine, L-threonine and L-isoleucine)(Miflin *et al.*, In: *Nitrogen Assimilation in Plants*, Hewitt *et al.*, (eds.), Academic Press, New York, 335 (1997); Bryan, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York, 403 (1980); Lea *et al.*, In: *The Chemistry and Biochemistry of Amino Acids*, Barrett *et al.*, (eds.), London, 5:197 (1985); Bryan, In: *The Biochemistry of Plants*, Miflin *et al.*, (eds.), Academic Press, San Diego, 16:161 (1990), all of which are herein incorporated by reference in their entirety).

The methionine-specific part of the aspartate pathway includes the following enzymes: aspartate kinase (EC 2.7.2.4), aspartate-semialdehyde dehydrogenase (EC 1.2.1.11), homoserine dehydrogenase (EC 1.1.1.3), homoserine kinase (EC 2.7.1.39), cystathionine γ -synthase (EC 4.2.99.9), cystathionine β -lyase (EC 4.4.1.8) and methionine synthase (EC 2.1.1.14).

Aspartate kinase catalyzes the first reaction of the pathway in which aspartate is converted to β -aspartyl phosphate. This enzyme has been isolated and characterized from plant sources including maize, barley, carrot, pea and soybean. These studies have revealed that there are multiple isoenzymes of aspartate kinase and the isoenzymes differ with respect to both

feedback inhibition sensitivity and expression profile (tissue and developmental stage).

Feedback inhibition is mediated by lysine and threonine. Transgenic plants which express an unregulated aspartate kinase have demonstrated increased flux through the aspartate pathway.

Pathway regulation is reported to be exerted, at least in part, via control of this enzyme's activity.

Aspartate semialdehyde dehydrogenase catalyses the second pathway reaction and converts β -aspartyl phosphate to aspartate semialdehyde via an NADPH-dependent reaction.

Gengenbach *et al.*, *Crop Science* 18:472-476 (1978), the entirety of which is herein incorporated by reference, report the isolation of aspartate semialdehyde dehydrogenase from maize suspension culture cells. These suspension cultures did not exhibit feedback inhibition of the enzyme in the presence of aspartate-derived amino acids, with the exception of methionine, for which some feedback sensitivity was observed. Aspartate semialdehyde dehydrogenase enzyme activity has been reported in maize shoot, maize root and maize kernel (Gengenbach *et al.*, *Crop Science* 18:472-476 (1978)).

Homoserine dehydrogenase catalyzes the next step of the pathway in which homoserine is generated from aspartate semialdehyde in a reaction requiring NADH or NADPH.

Homoserine dehydrogenase enzyme has been studied in higher plants and multiple isoenzyme forms have been reported (Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970); Gengenbach *et al.*, *Crop Science* 18:472-476 (1978); Dotson *et al.*, *Plant Physiology* 91:1602-1608 (1989); Dotson *et al.*, *Plant Physiology* 93:98-104 (1989); Azevedo *et al.*, *Phytochemistry* 31:3725-3730 (1992); Azevedo *et al.*, *Phytochemistry* 31:3731-3734 (1992); Brennecke *et al.*, *Phytochemistry* 41:707 (1996); Aarnes, *Plant Science Letters* 9:137-145 (1977); Bright *et al.*, *Biochemical Genetics* 200:229-243 (1982); Aruda *et al.*, *Plant Physiology*

76:442-446 (1984); Lea *et al.*, In: *Barley: Genetics, Molecular Biology and Biotechnology* Shewrey (ed.), CAB International, Oxford 181 (1992); Davies *et al.*, *Plant Science Letters* 9:323-332 (1977); Davies *et al.*, *Plant Physiology* 62:536-541 (1978); Matthews *et al.*, *Zeitschrift für Naturforschung, Section Bioscience* 34:1177-1185 (1979); Relton *et al.*, *Biochimica et Biophysica Acta* 953:48-60 (1988); Aarnes *et al.*, *Phytochemistry* 13:2717-2724 (1974); Lea *et al.*, *FEBS Letters* 98:165-168 (1979); Matthews *et al.*, *Canadian Journal of Botany* 57:299-304 (1979), all of which references are incorporated herein in their entirety). The isoenzymes have been found to differ with respect to sensitivity to threonine-mediated feedback inhibition, with both sensitive and insensitive forms being isolated from maize suspension cultures and seedlings (Miflin *et al.*, In: *Nitrogen Assimilation of Plants*, Hewitt *et al.*, (eds.), Academic Press, New York, 335 (1997); Bryan, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York, 5:403 (1980)).

There is evidence that plants also possess a bifunctional enzyme with both aspartate kinase and homoserine dehydrogenase activities (Lea *et al.*, In: *The Chemistry and Biochemistry of Amino Acids*, Barrett *et al.* (eds), London, 5:197 (1985), the entirety of which is herein incorporated by reference; Bryan, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York, 5:161 (1990), the entirety of which is herein incorporated by reference). Clones of these bifunctional enzymes have been isolated from *Arabidopsis thaliana* (Giovanelli *et al.*, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York 453 (1990), the entirety of which is herein incorporated by reference) carrot (Giovanelli *et al.*, *Plant Physiology* 90:1584-1599 (1989), the entirety of which is herein incorporated by reference), maize (Singh *et al.*, *Amino Acids* 7:165-168 (1994), the entirety of which is herein incorporated by reference) and soybean (Matthews *et al.*, In: *Biosynthesis and Molecular Regulation of Amino Acids in Plants*, p

294, Singh *et al.* (eds.), American Society of Plant Physiologists, Rockville, MD (1992), the entirety of which is herein incorporated by reference).

The next reported enzymatic step leading to methionine biosynthesis in higher plants is the final common reaction shared by other amino acid end products (threonine and isoleucine). The reaction is catalyzed by homoserine kinase and it generates *O*-phosphohomoserine from homoserine, with ATP serving as the phosphate donor. Exceptions are *Pisum sativum* and *Lathyrus sitivus* which synthesize *O*-acetylhomoserine and *O*-oxalylhomoserine, respectively (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997), the entirety of which is herein incorporated by reference). Enteric bacteria use *O*-succinylhomoserine, while several gram-positive bacteria, yeasts and fungi use *O*-acetylhomoserine (formed using homoserine *O*-acetyltransferase (EC 2.3.1.31) (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)). Homoserine kinase has been reported from multiple higher plant sources (Galili, *The Plant Cell* 7:899-906 (1995), the entirety of which is herein incorporated by reference; Rees *et al.*, *Biochemical Journal* 309:999-1107 (1995), the entirety of which is herein incorporated by reference; Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970), the entirety of which is herein incorporated by reference; Gengenbach *et al.*, *Crop Science* 18:472-476 (1978), Dotson *et al.*, *Plant Physiology* 91:1602-1608 (1989), the entirety of which is herein incorporated by reference; Dotson *et al.*, *Plant Physiology* 93:98-104 (1989), the entirety of which is herein incorporated by reference). Homoserine kinase isolated from barley and wheat has not been reported to exhibit aspartate-derived amino acid feedback inhibition (Gengenbach *et al.*, *Crop Science* 18:472-476 (1978); Dotson *et al.*, *Plant Physiology* 93:98-104 (1989)). It has been reported that homoserine kinase exhibits feedback regulation in the dicots, pea (Rees *et al.*, *Biochemical Journal* 309:999-1007 (1995), the entirety of which is herein

incorporated by reference) and radish (Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970)). Bacterial and yeast homologues have been reported (Azevedo *et al.*, *Phytochemistry* 31:3725-3730 (1992); Azevedo *et al.*, *Phytochemistry* 31:3731-3734 (1992); Brennecke *et al.*, *Phytochemistry* 41:707 (1996); Aarnes, *Plant Science Letters* 9:137-145 (1977)).

Sulfur, in yeast, is incorporated into *O*-acetylhomoserine resulting in homocysteine. This reaction is catalyzed by the *O*-acetylhomoserine sulfhydrylase (EC 4.2.99.10) (also known as *O*-acethomoserine (thiol)-lyase). *O*-acetylhomoserine sulfhydrylase has been reported to be a homotetramer with a molecular weight of 200,000. *O*-acetylhomoserine sulfhydrylase has also been reported to bind four molecules of pyridoxal phosphate (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

In higher plants, the sulfur atom from cysteine and the carbon backbone derived from aspartate used to synthesize methionine are reported to be catalyzed by pyridoxal 5'-phosphate (PLP) dependent enzymes (Ravanel *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 95:7805-7812 (1998), the entirety of which is herein incorporated by reference). The amino acid composition of the *O*-acetylhomoserine sulfhydrylase has also been reported to share sequence similarities to the *E. coli* cystathionine γ -synthase and cystathionine β -lyase and cystathionine γ -lyase from *Saccharomyces cerevisiae* and rats. All of these enzymes thus appear to belong to one protein family, whose members have evolved from an ancestral pyridoxal phosphate enzyme (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

In yeast, the synthesis of cysteine from homocysteine has been reported to require two successive steps, β addition and γ elimination. Cystathionine β -synthase (EC 4.2.1.22) has been reported to catalyze the first reaction where homocysteine and serine yield cystathionine. In *S.*

cervisiae, cystathionine β -synthase is encoded by STR4. STR4 encodes a polypeptide of 506 residues which shows extensive sequence similarity to its functional analog in rats. The rat analog has been reported to contain an additional amino-terminal extension of 60 residues. Moreover, the two enzymes have been reported to be closely related to the cyteine synthase from enteric bacteria and plants (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

Cystathionine γ -lyase (EC 4.4.1.1) catalyzes the γ cleavage of cystationine in yeast, the second reported step of the biosynthesis of cysteine from homocysteine. Cystathionine γ -lyase has been reported to have a molecular weight of about 194,000kd. In *S. cerviseae*, cystathionine γ -lyase is encoded by STR1. A mutation in the *S. cerviseae* cystathionine γ -lyase gene leads to a nutritional requirement for cysteine or glutathione. The yeast cystathionine γ -lyase belongs to a protein family which includes a functional analog in rats, a Met25p from yeast and cystathionin β -lyase and cystathionin γ -synthase from *E. coli* (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

Cystathionine γ -synthase (also known as *O*-succinylhomoserine (thio)-lyase, E.C. 4.2.99.9) catalyzes the first reported reaction which is unique to methionine biosynthesis, thereby committing aspartate pathway flux toward this amino acid. In this reaction, *O*-phosphohomoserine and cysteine serve as substrates for the production of cystathionine. Cystathionine γ -synthase has not been reported to be regulated by aspartate-derived amino acids feedback inhibition (Bright *et al.*, *Biochemical Genetics* 20:229-243 (1982); Arruda *et al.*, *Plant Physiology* 76:442-446 (1984)). Cystathionine γ -synthase has however, been reported to be sensitive to product inhibition by orthophosphate (Lea *et al.*, *Barley: Genetics, Molecular*

Biology and Biotechnology, Shewrey (ed.), CAB International, Oxford, 181 (1992); Davies *et al.*, *Plant Science Letters* 9:323-332 (1977)). Cloned cystathionine γ -synthase have been reported from *Arabidopsis thaliana* (Davies *et al.*, *Plant Physiology* 62:536-541 (1978)). It has been reported that methionine levels are modulated via regulation of cystathionine-synthase (Matthews *et al.*, *Zeitschrift für Naturforschung, Section Bioscience* 34:1177-1185 (1979-2724 (1974); Lea *et al.*, *FEBS Letters* 98:165 (1979), all of which references are incorporated herein in their entirety).

Cystathionine β -lyase catalyzes the next reaction in the biosynthesis of methionine. This reaction generates homocysteine, pyruvate and ammonia from the enzymatic decomposition of cystathionine. Evidence for isoenzymes which differ with respect to cellular localization have been reported for barley (Matthews *et al.*, *Canadian Journal of Botany* 57:299-304 (1979)) and spinach (Rognes *et al.*, *Nature* 287:357-359 (1980), the entirety of which is herein incorporated by reference).

De novo synthesis of methionine from homocysteine uses a methyl group which originates from single-carbon metabolism. In this metabolism, derivatives of tetrahydrofolate transfer one-carbon groups at the oxidation levels of methanol, formaldehyde and formate to acceptor molecules. Single-carbon derivatives of tetrahydrofolate are required for the biosynthesis of methionine, purine nucleotides and thymidylate as well as for the synthesis of N-formylmethionine in the mitochondrion. *S. cerevisiae* possesses two complete sets of folate interconversion enzymes, one located in the cytosol (methionyl-tRNA synthetase, EC 6.1.1.10) and the other located in the mitochondrion (methionyl t-RNA synthetase, EC 6.1.1.10) (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)) and in plants including the

chloroplast (Menand *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)*, 95:11014-11019 (1998), the entirety of which is herein incorporated by reference).

Methionine synthase generates methionine from homocysteine by a methylation reaction and thus represents the final step of the methionine biosynthetic pathway. Methionine synthase is also sometimes referred to as 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase. N-methyltetrahydrofolate serves as the methyl donor in this reaction, which occurs in the absence of cobalamin (Giovanelli *et al.*, *Plant Physiology* 90:1577-1583 (1989), the entirety of which is herein incorporated by reference; Green *et al.*, *Crop Science* 14:827-830 (1974), the entirety of which is herein incorporated by reference).

II. METHIONINE DEGRADATION PATHWAY

Plants contain a pathway for the degradation of L-methionine. This degradation pathway includes the following enzymes: methionine adenosyltransferase (EC 2.5.1.6), methionine S-methyltransferase (EC 2.1.1.12), adenosylmethionine hydrolase (EC 3.3.1.2), homocysteine S-methyltransferase (EC 2.1.1.10) and S-adenosyl-methionine decarboxylase (EC 4.1.1.50).

The reported first step in the catabolism of methionine is the ATP-dependent conversion to S-adenosylmethionine (AdoMet), which is catalyzed by the enzyme methionine adenosyltransferase, also known as S-adenosylmethionine synthetase. Methionine adenosyltransferase enzyme has been characterized from several plant sources (Aarnes, *Plant Science Letters* 10:381 (1977), the entirety of which is herein incorporated by reference; Mathur *et al.*, *Biochimica and Biophysica Acta* 1078:161-170 (1991), the entirety of which is herein incorporated by reference; Kim *et al.*, *Journal of Biochemical and Molecular Biology* 28:100 (1995), the entirety of which is herein incorporated by reference) and nucleic acid molecules (genomic and cDNA) have also been obtained from a variety of sources (Izhaki *et al.*, *Plant*

Physiology 108:841-842 (1995), the entirety of which is herein incorporated by reference; Espartero *et al.*, *Molecular Biology Plant* 25:217-237 (1994), the entirety of which is herein incorporated by reference). Regulation of methionine adenosyltransferase activity has been observed for the enzyme from *Glycine max* (soybean). In *Glycine max*, methionine adenosyltransferase was reportedly inhibited by S-adenosylmethionine (Kim *et al.*, *Journal of Biochemical and Molecular Biology* 28:100 (1995). Studies have also reported that the levels of methionine adenosyltransferase appear to fluctuate in response to hormonal or environmental conditions such as gibberellic acid (Mathur *et al.*, *Biochimica and Biophysica Acta* 1162:289-290 (1993), the entirety of which is herein incorporated by reference; Mathur *et al.*, *Biochimica and Biophysica Acta* 1137:338-348 (1992), the entirety of which is herein incorporated by reference), salt stress (Espartero *et al.*, *Molecular Biology Plant* 25:217-227 (1994) the entirety of which is herein incorporated by reference) and wounding (Kim *et al.*, *Plant Cell Reports* 13:340 (1994), the entirety of which is herein incorporated by reference). It has also been reported that methionine adenosyltransferase may play a role in the lignification process (Peleman *et al.*, *Plant Cell* 1:81 (1989), the entirety of which is herein incorporated by reference).

AdoMet is further catabolized by several enzymes and has been reported to serve a variety of metabolic functions including that of a methyl donor (Cossins, *The Biochemistry of Plants* 11:317 Devis (ed.), Academic Press, San Diego (1987), the entirety of which is herein incorporated by reference) that of a precursor for polyamine biosynthesis (Tiburico *et al.*, *The Biochemistry of Plants* 16:283 (1990), the entirety of which is herein incorporated by reference) and that of a precursor for ethylene biosynthesis (Kende, *Plant Physiology* 91:1-4 (1989), the entirety of which is herein incorporated by reference; Flurh *et al.*, *Critical Review of Plant*

Science 15:479 (1996), the entirety of which is herein incorporated by reference). In each case, enzymes are present to regenerate methionine from the sulfur-containing backbone resulting in no net loss of methionine.

An enzyme involved in AdoMet catabolism is adenosylmethionine hydrolase (EC 3.3.1.2) which converts AdoMet to methylthioadenosine and L-homoserine. L-homoserine is further metabolized during the biosynthesis of polyamines and ethylene and methylthioadenosine is recycled to methionine. In yeast, a form of adenosylmethionine hydrolase (EC 3.1.1.1) has been reported (<http://www.ncbi.nlm.nih.gov/htbin-post/Entrez/query> (1998)).

Another enzyme for which AdoMet is a substrate for is homocysteine S-methyltransferase. Homocysteine S-methyltransferase catalyzes the combination of AdoMet, with L-homocysteine to produce both S-adenosyl-L-homocysteine and L-methionine. Another enzyme has been described which generates S-adenosyl-L-homocysteine from AdoMet. This enzyme is called methionine S-methyltransferase and it catalyzes the reaction in which S-adenosyl-L-homocysteine reacts with L-methionine to generate S-adenosyl-L-homocysteine and S-methyl-L-methionine. AdoMet can also be decarboxylated by adenosyl methionine decarboxylase, which generates (5-deoxy-5-adenosyl) (3-aminopropyl) methylsulfonium salt.

III. EXPRESSED SEQUENCE TAG NUCLEIC ACID MOLECULES

Expressed sequence tags, or ESTs are randomly sequenced members of a cDNA library (or complementary DNA)(McCombie *et al.*, *Nature Genetics* 1:124-130 (1992); Kurata *et al.*, *Nature Genetics* 8:365-372 (1994); Okubo *et al.*, *Nature Genetics* 2:173-179 (1992), all of which references are incorporated herein in their entirety). The randomly selected clones comprise insets that can represent a copy of up to the full length of a mRNA transcript.

Using conventional methodologies, cDNA libraries can be constructed from the mRNA (messenger RNA) of a given tissue or organism using poly dT primers and reverse transcriptase (Efstratiadis *et al.*, *Cell* 7:279-3680 (1976), the entirety of which is herein incorporated by reference; Higuchi *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 73:3146-3150 (1976), the entirety of which is herein incorporated by reference; Maniatis *et al.*, *Cell* 8:163-182 (1976) the entirety of which is herein incorporated by reference; Land *et al.*, *Nucleic Acids Res.* 9:2251-2266 (1981), the entirety of which is herein incorporated by reference; Okayama *et al.*, *Mol. Cell. Biol.* 2:161-170 (1982), the entirety of which is herein incorporated by reference; Gubler *et al.*, *Gene* 25:263-269 (1983), the entirety of which is herein incorporated by reference).

Several methods may be employed to obtain full-length cDNA constructs. For example, terminal transferase can be used to add homopolymeric tails of dC residues to the free 3' hydroxyl groups (Land *et al.*, *Nucleic Acids Res.* 9:2251-2266 (1981), the entirety of which is herein incorporated by reference). This tail can then be hybridized by a poly dG oligo which can act as a primer for the synthesis of full length second strand cDNA. Okayama and Berg, *Mol. Cell. Biol.* 2:161-170 (1982), the entirety of which is herein incorporated by reference, report a method for obtaining full length cDNA constructs. This method has been simplified by using synthetic primer-adapters that have both homopolymeric tails for priming the synthesis of the first and second strands and restriction sites for cloning into plasmids (Coleclough *et al.*, *Gene* 34:305-314 (1985), the entirety of which is herein incorporated by reference) and bacteriophage vectors (Krawinkel *et al.*, *Nucleic Acids Res.* 14:1913 (1986), the entirety of which is herein incorporated by reference; Han *et al.*, *Nucleic Acids Res.* 15:6304 (1987), the entirety of which is herein incorporated by reference).

These strategies have been coupled with additional strategies for isolating rare mRNA populations. For example, a typical mammalian cell contains between 10,000 and 30,000 different mRNA sequences (Davidson, *Gene Activity in Early Development*, 2nd ed., Academic Press, New York (1976), the entirety of which is herein incorporated by reference). The number of clones required to achieve a given probability that a low-abundance mRNA will be present in a cDNA library is $N = (\ln(1-P))/(\ln(1-1/n))$ where N is the number of clones required, P is the probability desired and 1/n is the fractional proportion of the total mRNA that is represented by a single rare mRNA (Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, 2nd ed., Cold Spring Harbor Laboratory Press (1989), the entirety of which is herein incorporated by reference).

A method to enrich preparations of mRNA for sequences of interest is to fractionate by size. One such method is to fractionate by electrophoresis through an agarose gel (Pennica *et al.*, *Nature* 301:214-221 (1983), the entirety of which is herein incorporated by reference). Another such method employs sucrose gradient centrifugation in the presence of an agent, such as methylmercuric hydroxide, that denatures secondary structure in RNA (Schweinfest *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 79:4997-5000 (1982), the entirety of which is herein incorporated by reference).

A frequently adopted method is to construct equalized or normalized cDNA libraries (Ko, *Nucleic Acids Res.* 18:5705-5711 (1990), the entirety of which is herein incorporated by reference; Patanjali *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:1943-1947 (1991), the entirety of which is herein incorporated by reference). Typically, the cDNA population is normalized by subtractive hybridization (Schmid *et al.*, *J. Neurochem.* 48:307-312 (1987), the entirety of which is herein incorporated by reference; Fargnoli *et al.*, *Anal. Biochem.* 187:364-373 (1990), the

entirety of which is herein incorporated by reference; Travis *et al.*, *Proc. Natl. Acad. Sci (U.S.A.)* 85:1696-1700 (1988), the entirety of which is herein incorporated by reference; Kato, *Eur. J. Neurosci.* 2:704-711 (1990); and Schweinfest *et al.*, *Genet. Anal. Tech. Appl.* 7:64-70 (1990), the entirety of which is herein incorporated by reference). Subtraction represents another method for reducing the population of certain sequences in the cDNA library (Swaroop *et al.*, *Nucleic Acids Res.* 19:1954 (1991), the entirety of which is herein incorporated by reference).

ESTs can be sequenced by a number of methods. Two basic methods may be used for DNA sequencing, the chain termination method of Sanger *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 74:5463-5467 (1977), the entirety of which is herein incorporated by reference and the chemical degradation method of Maxam and Gilbert, *Proc. Nat. Acad. Sci. (U.S.A.)* 74:560-564 (1977), the entirety of which is herein incorporated by reference. Automation and advances in technology such as the replacement of radioisotopes with fluorescence-based sequencing have reduced the effort required to sequence DNA (Craxton, *Methods* 2:20-26 (1991), the entirety of which is herein incorporated by reference; Ju *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 92:4347-4351 (1995), the entirety of which is herein incorporated by reference; Tabor and Richardson, *Proc. Natl. Acad. Sci. (U.S.A.)* 92:6339-6343 (1995), the entirety of which is herein incorporated by reference). Automated sequencers are available from, for example, Pharmacia Biotech, Inc., Piscataway, New Jersey (Pharmacia ALF), LI-COR, Inc., Lincoln, Nebraska (LI-COR 4,000) and Millipore, Bedford, Massachusetts (Millipore BaseStation).

In addition, advances in capillary gel electrophoresis have also reduced the effort required to sequence DNA and such advances provide a rapid high resolution approach for sequencing DNA samples (Swerdlow and Gesteland, *Nucleic Acids Res.* 18:1415-1419 (1990); Smith, *Nature* 349:812-813 (1991); Luckey *et al.*, *Methods Enzymol.* 218:154-172 (1993); Lu *et al.*, *J.*

Chromatog. A. 680:497-501 (1994); Carson *et al.*, *Anal. Chem.* 65:3219-3226 (1993); Huang *et al.*, *Anal. Chem.* 64:2149-2154 (1992); Kheterpal *et al.*, *Electrophoresis* 17:1852-1859 (1996); Quesada and Zhang, *Electrophoresis* 17:1841-1851 (1996); Baba, *Yakugaku Zasshi* 117:265-281 (1997), all of which are herein incorporated by reference in their entirety).

ESTs longer than 150 nucleotides have been found to be useful for similarity searches and mapping (Adams *et al.*, *Science* 252:1651-1656 (1991), herein incorporated by reference).

ESTs, which can represent copies of up to the full length transcript, may be partially or completely sequenced. Between 150-450 nucleotides of sequence information is usually generated as this is the length of sequence information that is routinely and reliably produced using single run sequence data. Typically, only single run sequence data is obtained from the cDNA library (Adams *et al.*, *Science* 252:1651-1656 (1991). Automated single run sequencing typically results in an approximately 2-3% error or base ambiguity rate (Boguski *et al.*, *Nature Genetics* 4:332-333 (1993), the entirety of which is herein incorporated by reference).

EST databases have been constructed or partially constructed from, for example, *C. elegans* (McCombie *et al.*, *Nature Genetics* 1:124-131 (1992)), human liver cell line HepG2 (Okubo *et al.*, *Nature Genetics* 2:173-179 (1992)), human brain RNA (Adams *et al.*, *Science* 252:1651-1656 (1991); Adams *et al.*, *Nature* 355:632-635 (1992)), *Arabidopsis*, (Newman *et al.*, *Plant Physiol.* 106:1241-1255 (1994)); and rice (Kurata *et al.*, *Nature Genetics* 8:365-372 (1994)).

IV. SEQUENCE COMPARISONS

A characteristic feature of a DNA sequence is that it can be compared with other DNA sequences. Sequence comparisons can be undertaken by determining the similarity of the test or query sequence with sequences in publicly available or proprietary databases ("similarity

analysis”) or by searching for certain motifs (“intrinsic sequence analysis”)(e.g. *cis* elements)(Coulson, *Trends in Biotechnology* 12:76-80 (1994), the entirety of which is herein incorporated by reference); Birren *et al.*, *Genome Analysis 1*: Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York 543-559 (1997), the entirety of which is herein incorporated by reference).

Similarity analysis includes database search and alignment. Examples of public databases include the DNA Database of Japan (DDBJ)(<http://www.ddbj.nig.ac.jp/>); Genebank (<http://www.ncbi.nlm.nih.gov/Web/Search/Index.html>); and the European Molecular Biology Laboratory Nucleic Acid Sequence Database (EMBL) (http://www.ebi.ac.uk/ebi_docs/embl_db/embl-db.html). Other appropriate databases include dbEST (<http://www.ncbi.nlm.nih.gov/dbEST/index.html>), SwissProt (http://www.ebi.ac.uk/ebi_docs/swisprot_db/swisshome.html), PIR (<http://www-nbrt.georgetown.edu/pir/>) and The Institute for Genome Research (<http://www.tigr.org/tdb/tdb.html>)

A number of different search algorithms have been developed, one example of which are the suite of programs referred to as BLAST programs. There are five implementations of BLAST, three designed for nucleotide sequences queries (BLASTN, BLASTX and TBLASTX) and two designed for protein sequence queries (BLASTP and TBLASTN) (Coulson, *Trends in Biotechnology* 12:76-80 (1994); Birren *et al.*, *Genome Analysis 1*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York 543-559 (1997)).

BLASTN takes a nucleotide sequence (the query sequence) and its reverse complement and searches them against a nucleotide sequence database. BLASTN was designed for speed, not maximum sensitivity and may not find distantly related coding sequences. BLASTX takes a

nucleotide sequence, translates it in three forward reading frames and three reverse complement reading frames and then compares the six translations against a protein sequence database.

BLASTX is useful for sensitive analysis of preliminary (single-pass) sequence data and is tolerant of sequencing errors (Gish and States, *Nature Genetics* 3:266-272 (1993), the entirety of which is herein incorporated by reference). BLASTN and BLASTX may be used in concert for analyzing EST data (Coulson, *Trends in Biotechnology* 12:76-80 (1994); Birren *et al.*, *Genome Analysis* 1:543-559 (1997)).

Given a coding nucleotide sequence and the protein it encodes, it is often preferable to use the protein as the query sequence to search a database because of the greatly increased sensitivity to detect more subtle relationships. This is due to the larger alphabet of proteins (20 amino acids) compared with the alphabet of nucleic acid sequences (4 bases), where it is far easier to obtain a match by chance. In addition, with nucleotide alignments, only a match (positive score) or a mismatch (negative score) is obtained, but with proteins, the presence of conservative amino acid substitutions can be taken into account. Here, a mismatch may yield a positive score if the non-identical residue has physical/chemical properties similar to the one it replaced. Various scoring matrices are used to supply the substitution scores of all possible amino acid pairs. A general purpose scoring system is the BLOSUM62 matrix (Henikoff and Henikoff, *Proteins* 17:49-61 (1993), the entirety of which is herein incorporated by reference), which is currently the default choice for BLAST programs. BLOSUM62 is tailored for alignments of moderately diverged sequences and thus may not yield the best results under all conditions. Altschul, *J. Mol. Biol.* 36:290-300 (1993), the entirety of which is herein incorporated by reference, describes a combination of three matrices to cover all contingencies. This may improve sensitivity, but at the expense of slower searches. In practice, a single

BLOSUM62 matrix is often used but others (PAM40 and PAM250) may be attempted when additional analysis is necessary. Low PAM matrices are directed at detecting very strong but localized sequence similarities, whereas high PAM matrices are directed at detecting long but weak alignments between very distantly related sequences.

Homologues in other organisms are available that can be used for comparative sequence analysis. Multiple alignments are performed to study similarities and differences in a group of related sequences. CLUSTAL W is a multiple sequence alignment package that performs progressive multiple sequence alignments based on the method of Feng and Doolittle, *J. Mol. Evol.* 25:351-360 (1987), the entirety of which is herein incorporated by reference. Each pair of sequences is aligned and the distance between each pair is calculated; from this distance matrix, a guide tree is calculated and all of the sequences are progressively aligned based on this tree. A feature of the program is its sensitivity to the effect of gaps on the alignment; gap penalties are varied to encourage the insertion of gaps in probable loop regions instead of in the middle of structured regions. Users can specify gap penalties, choose between a number of scoring matrices, or supply their own scoring matrix for both pairwise alignments and multiple alignments. CLUSTAL W for UNIX and VMS systems is available at: <ftp://ebi.ac.uk>. Another program is MACAW (Schuler *et al.*, *Proteins Struct. Func. Genet.* 9:180-190 (1991), the entirety of which is herein incorporated by reference, for which both Macintosh and Microsoft Windows versions are available. MACAW uses a graphical interface, provides a choice of several alignment algorithms and is available by anonymous ftp at: [ncbi.nlm.nih.gov](ftp://ncbi.nlm.nih.gov) (directory/pub/macaw).

Sequence motifs are derived from multiple alignments and can be used to examine individual sequences or an entire database for subtle patterns. With motifs, it is sometimes

possible to detect distant relationships that may not be demonstrable based on comparisons of primary sequences alone. Currently, the largest collection of sequence motifs in the world is PROSITE (Bairoch and Bucher, *Nucleic Acid Research* 22:3583-3589 (1994), the entirety of which is herein incorporated by reference). PROSITE may be accessed via either the ExPASy server on the World Wide Web or anonymous ftp site. Many commercial sequence analysis packages also provide search programs that use PROSITE data.

A resource for searching protein motifs is the BLOCKS E-mail server developed by Henikoff, *Trends Biochem Sci.* 18:267-268 (1993), the entirety of which is herein incorporated by reference; Henikoff and Henikoff, *Nucleic Acid Research* 19:6565-6572 (1991), the entirety of which is herein incorporated by reference; Henikoff and Henikoff, *Proteins* 17:49-61 (1993). BLOCKS searches a protein or nucleotide sequence against a database of protein motifs or “blocks.” Blocks are defined as short, ungapped multiple alignments that represent highly conserved protein patterns. The blocks themselves are derived from entries in PROSITE as well as other sources. Either a protein query or a nucleotide query can be submitted to the BLOCKS server; if a nucleotide sequence is submitted, the sequence is translated in all six reading frames and motifs are sought for these conceptual translations. Once the search is completed, the server will return a ranked list of significant matches, along with an alignment of the query sequence to the matched BLOCKS entries.

Conserved protein domains can be represented by two-dimensional matrices, which measure either the frequency or probability of the occurrences of each amino acid residue and deletions or insertions in each position of the domain. This type of model, when used to search against protein databases, is sensitive and usually yields more accurate results than simple motif searches. Two popular implementations of this approach are profile searches such as GCG

program ProfileSearch and Hidden Markov Models (HMMs)(Krough *et al.*, *J. Mol. Biol.* 235:1501-1531, (1994); Eddy, *Current Opinion in Structural Biology* 6:361-365, (1996), both of which are herein incorporated by reference in their entirety). In both cases, a large number of common protein domains have been converted into profiles, as present in the PROSITE library, or HMM models, as in the Pfam protein domain library (Sonnhammer *et al.*, *Proteins* 28:405-420 (1997), the entirety of which is herein incorporated by reference). Pfam contains more than 500 HMM models for enzymes, transcription factors, signal transduction molecules and structural proteins. Protein databases can be queried with these profiles or HMM models, which will identify proteins containing the domain of interest. For example, HMMSW or HMMFS, two programs in a public domain package called HMMER (Sonnhammer *et al.*, *Proteins* 28:405-420 (1997)) can be used.

PROSITE and BLOCKS represent collected families of protein motifs. Thus, searching these databases entails submitting a single sequence to determine whether or not that sequence is similar to the members of an established family. Programs working in the opposite direction compare a collection of sequences with individual entries in the protein databases. An example of such a program is the Motif Search Tool, or MoST (Tatusov *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:12091-12095 (1994), the entirety of which is herein incorporated by reference). On the basis of an aligned set of input sequences, a weight matrix is calculated by using one of four methods (selected by the user). A weight matrix is simply a representation, position by position of how likely a particular amino acid will appear. The calculated weight matrix is then used to search the databases. To increase sensitivity, newly found sequences are added to the original data set, the weight matrix is recalculated and the search is performed again. This procedure continues until no new sequences are found.

SUMMARY OF THE INVENTION

The present invention provides a substantially purified nucleic acid molecule that encodes a maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of: (a) methionine adenosyltransferase, (b) S-adenosyl-methionine decarboxylase, (c) aspartate kinase, (d) aspartate-semialdehyde dehydrogenase, (e) cystathionine gamma-synthase, (f) cystathionine beta-lyase, and (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase.

The present invention also provides a substantially purified nucleic acid molecule that encodes a plant methionine pathway enzyme or fragment thereof, wherein the nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a

maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof.

A substantially purified maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of (a) methionine adenosyltransferase or fragment thereof; (b) S-adenosyl-methionine decarboxylase or fragment thereof; (c) aspartate kinase or fragment thereof; (d) aspartate-semialdehyde dehydrogenase or fragment thereof; (e) cystathionine gamma-synthase or fragment thereof; (f) cystathionine beta-lysase or fragment thereof; and (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof.

The present invention also provides a substantially purified maize or soybean methionine pathway protein or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified maize or soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a

complement of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.

The present invention also provides a substantially purified maize or soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.

The present invention also provides a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified maize or soybean cystathionine β -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified maize or soybean cystathionine β -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected

from the group consisting of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified maize or adenosylhomocysteinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified maize or adenosylhomocysteinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified maize or cystathionine β -synthase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified maize or cystathionine β -synthase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified maize or cystathionine γ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or cystathionine γ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a substantially purified maize or *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a purified antibody or fragment thereof which is capable of specifically binding to a specific maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme or fragment thereof is encoded by a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ

ID NO: 2623 or a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean aspartate kinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or a substantially purified maize or soybean enzyme aspartate-semialdehyde dehydrogenase or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine β -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665 or a substantially purified maize or soybean cystathionine β -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic

acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199 or a substantially purified maize or soybean adenosylhomocysteinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine β -synthase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1631 through SEQ ID NO: 1632 or a substantially purified maize or soybean cystathionine β -synthase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204 or a substantially purified maize or soybean cystathionine γ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean *O*-acetylhomoserine enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 3200 through SEQ ID NO: 3202 or a substantially purified maize or soybean *O*-acetylhomoserine enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; (B) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of (a) a nucleic acid sequence which encodes for methionine adenosyltransferase or fragment thereof; (b) a nucleic acid sequence which encodes for S-adenosyl-methionine decarboxylase or fragment thereof; (c) a nucleic acid

sequence which encodes for aspartate kinase or fragment thereof; (d) a nucleic acid sequence which encodes for aspartate-semialdehyde dehydrogenase or fragment thereof; (e) a nucleic acid sequence which encodes for cystathionine gamma-synthase or a fragment thereof; (f) a nucleic acid sequence which encodes for cystathionine beta-lyase or a fragment thereof; (g) a nucleic acid sequence which encodes for 5-methyltetrahydropteroyl- tri glutamate-homocysteine-S-methyltransferase or a fragment thereof; and (h) a nucleic acid sequence which is complementary to any of the nucleic acid sequences of (a) through (g); and (C) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a structural nucleic acid molecule, wherein the structural nucleic acid molecule encodes a plant methionine pathway enzyme or fragment thereof, the structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; which is linked to (C) a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a structural nucleic acid molecule, wherein the structural nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or

fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; which is linked to (C) a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a transcribed nucleic acid molecule with a transcribed strand and a non-transcribed strand, wherein the transcribed strand is complementary to a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; which is

linked to (C) a 3' non-translated sequence that functions in plant cells to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to: (B) a transcribed nucleic acid molecule with a transcribed strand and a non-transcribed strand, wherein a transcribed mRNA of the transcribed strand is complementary to an endogenous mRNA molecule having a nucleic acid sequence selected from the group consisting of an endogenous mRNA molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and an endogenous mRNA molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; which is linked to (C) a 3'

non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a method for determining a level or pattern in a plant cell of an enzyme in a plant metabolic pathway comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence of SEQ ID NO: 1 through SEQ ID NO: 3204 or compliments thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said enzyme; (B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and (C) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said enzyme in said plant metabolic pathway.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complement thereof or fragment of either, with a complementary nucleic acid molecule obtained from the plant cell or plant tissue, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue permits the detection of the

plant methionine pathway enzyme; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) detecting the level or pattern of the complementary nucleic acid, wherein the detection of the complementary nucleic acid is predictive of the level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a

soybean cytsathionine γ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, with a complementary nucleic acid molecule obtained from the plant cell or plant tissue, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue permits the detection of the plant methionine pathway enzyme; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) detecting the level or pattern of the complementary nucleic acid, wherein the detection of the complementary nucleic acid is predictive of the level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue under evaluation which comprises assaying the concentration of a molecule, whose concentration is dependent upon the expression of a gene, the gene specifically hybridizes to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof, in comparison to the concentration of that molecule present in a reference plant cell or a reference plant tissue with a known level or pattern of the plant methionine pathway enzyme, wherein the assayed concentration of the molecule is compared to the assayed concentration of the molecule in the reference plant cell or reference plant tissue with the known level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue under evaluation which comprises assaying the concentration of a molecule, whose concentration is dependent upon the expression

of a gene, the gene specifically hybridizes to a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or complement thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof, in comparison to the concentration of that molecule present in a reference plant cell or a reference plant tissue with a known level or pattern of the plant methionine pathway enzyme, wherein the assayed concentration of the molecule is compared to the assayed concentration of the molecule in the reference plant cell or the reference plant tissue with the known level or pattern of the plant methionine pathway enzyme.

A method of determining a mutation in a plant whose presence is predictive of a mutation affecting a level or pattern of a protein comprising the steps: (A) incubating, under conditions

permitting nucleic acid hybridization, a marker nucleic acid, said marker nucleic acid selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having a nucleic acid sequence selected from the group of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from said plant, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting said level or pattern of said plant methionine pathway enzyme in said plant; (B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant; and (C) detecting the presence of said polymorphism, wherein the detection of said polymorphism is predictive of said mutation.

The present invention also provides a method for determining a mutation in a plant whose presence is predictive of a mutation affecting the level or pattern of a plant methionine pathway enzyme comprising the steps: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that is linked to a gene, the gene specifically hybridizes to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from the plant, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting the level or pattern of the plant methionine pathway enzyme in the plant; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule

obtained from the plant; and (C) detecting the presence of the polymorphism, wherein the detection of the polymorphism is predictive of the mutation.

The present invention also provides a method for determining a mutation in a plant whose presence is predictive of a mutation affecting the level or pattern of a plant methionine pathway enzyme comprising the steps: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that is linked to a gene, the gene specifically hybridizes to a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or complement thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof and a complementary nucleic acid molecule obtained from the plant,

wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting the level or pattern of the plant methionine pathway enzyme in the plant; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant; and (C) detecting the presence of the polymorphism, wherein the detection of the polymorphism is predictive of the mutation.

A method of producing a plant containing an overexpressed protein comprising: (A) transforming said plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein said promoter region is linked to a structural region, wherein said structural region has a nucleic acid sequence selected from group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 wherein said structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein said functional nucleic acid molecule results in overexpression of the protein; and (B) growing said transformed plant.

The present invention also provides a method of producing a plant containing an overexpressed plant methionine enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause

termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in overexpression of the plant methionine pathway enzyme; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing an overexpressed plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; wherein the structural region is

linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in overexpression of the plant methionine pathway enzyme protein; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing reduced levels of a plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in co-suppression of the plant methionine pathway enzyme protein; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing reduced levels of a plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a

nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in co-suppression of the plant methionine pathway enzyme; and (B) growing the transformed plant.

The present invention also provides a method for reducing expression of a plant methionine pathway enzyme in a plant comprising: (A) transforming the plant with a nucleic acid molecule, the nucleic acid molecule having an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule, wherein the exogenous promoter region is linked to a transcribed nucleic acid molecule having a transcribed strand and a non-transcribed strand, wherein the transcribed strand is complementary to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either and the transcribed strand is complementary

to an endogenous mRNA molecule; and wherein the transcribed nucleic acid molecule is linked to a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and (B) growing the transformed plant.

The present invention also provides a method for reducing expression of a plant methionine pathway enzyme in a plant comprising: (A) transforming the plant with a nucleic acid molecule, the nucleic acid molecule having an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule, wherein the exogenous promoter region is linked to a transcribed nucleic acid molecule having a transcribed strand and a non-transcribed strand, wherein a transcribed mRNA of the transcribed strand is complementary to a nucleic acid molecule selected from the group consisting of an endogenous mRNA molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment

thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and an endogenous mRNA molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; and wherein the transcribed nucleic acid molecule is linked to a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and (B) growing the transformed plant.

The present invention also provides a method of determining an association between a polymorphism and a plant trait comprising: (A) hybridizing a nucleic acid molecule specific for the polymorphism to genetic material of a plant, wherein the nucleic acid molecule has a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragment thereof; and (B) calculating the degree of association between the polymorphism and the plant trait.

The present invention also provides a method of determining an association between a polymorphism and a plant trait comprising: (A) hybridizing a nucleic acid molecule specific for the polymorphism to genetic material of a plant, wherein the nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme complement thereof or fragment of either, a

nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either; a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine γ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either and (B) calculating the degree of association between the polymorphism and the plant trait.

The present invention also provides a method of isolating a nucleic acid that encodes a plant methionine pathway enzyme or fragment thereof comprising: (A) incubating under conditions permitting nucleic acid hybridization, a first nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragment of either with a complementary second nucleic acid molecule obtained from a plant cell or plant tissue; (B) permitting hybridization between the first nucleic acid molecule and the second nucleic acid molecule obtained from the plant cell or plant tissue; and (C) isolating the second nucleic acid molecule.

The present invention also provides a method of isolating a nucleic acid molecule that encodes a plant methionine pathway enzyme or fragment thereof comprising: (A) incubating under conditions permitting nucleic acid hybridization, a first nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean

methionine adenosyltransferase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine γ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, with a complementary second nucleic acid molecule obtained from a plant cell or plant tissue; (B) permitting hybridization between the plant methionine pathway nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) isolating the second nucleic acid molecule.

DETAILED DESCRIPTION OF THE INVENTION

Definitions and Agents of the Present Invention

Definitions:

As used herein, a methionine pathway enzyme is any enzyme that is associated with the synthesis or degradation of methionine.

As used herein, a methionine synthesis enzyme is any enzyme that is associated with the synthesis of methionine.

As used herein, a methionine degradation enzyme is any enzyme that is associated with the degradation of methionine.

As used herein, methionine adenosyltransferase is any enzyme that catalyzes the conversion of methionine to S-adenosylmethionine.

As used herein, S-adenosylmethionine decarboxylase is any enzyme that catalyzes the reaction that converts S-adenosylmethionine to (5-deoxy-5-adenosyl)(3-aminopropyl) methylsulfonium salt.

As used herein, aspartate kinase is any enzyme that catalyzes the conversion of aspartate to β -aspartyl phosphate.

As used herein, aspartate semialdehyde dehydrogenase is any enzyme that catalyzes the conversion of β -aspartyl phosphate to aspartate-semialdehyde via an NADPH-dependent reaction.

As used herein, *O*-succinylhomoserine (thiol)-lyase refers to any enzyme that catalyzes the conversion of *O*-phosphohomoserine to and cysteine to cystathionine.

As used herein, cystathionine β -lyase is any enzyme that catalyzes the conversion of cystathionine to homocysteine, pyruvate and ammonia.

As used herein, 5-methyltetrahydropterolytriglutamate-homocysteine S-methyltransferase refers to any enzyme which catalyzes the conversion of homocysteine via methylation to methionine.

As used herein, adenosylhomocysteinase refers to any enzyme that catalyzes the ATP-dependent conversion of S-adenosylmethionine (AdoMet) to methylthioadenosine and L-homoserine.

As used herein, cystathionine β -synthase refers to any enzyme that catalyzes the conversion of homocysteine and serine to cystathionine.

As used herein, cystathionine γ -lyase refers to any enzyme that catalyzes the γ cleavage of cystathionine.

As used herein, *O*-acetylhomoserine (thiol)-lyase refers to any enzyme that catalyzes the conversion of *O*-acetylhomoserine and sulfur to homocysteine.

Agents

(a) Nucleic Acid Molecules

Agents of the present invention include plant nucleic acid molecules and more specifically include maize and soybean nucleic acid molecules and more specifically include nucleic acid molecules of the maize genotypes B73 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), B73 x Mo17 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), DK604 (Dekalb Genetics, Dekalb, Illinois U.S.A.), H99 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), RX601 (Asgrow Seed Company, Des Moines, Iowa), Mo17 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), and soybean types Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa), C1944 (United States Department of Agriculture (USDA) Soybean Germplasm Collection, Urbana, Illinois U.S.A.), Cristalina (USDA Soybean Germplasm Collection, Urbana,

Illinois U.S.A.), FT108 (Monsoy, Brazil), Hartwig (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), BW211S Null (Tohoku University, Morioka, Japan), PI507354 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), Asgrow A4922 (Asgrow Seed Company, Des Moines, Iowa U.S.A.), PI227687 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), PI229358 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and Asgrow A3237 (Asgrow Seed Company, Des Moines, Iowa U.S.A.).

A subset of the nucleic acid molecules of the present invention includes nucleic acid molecules that are marker molecules. Another subset of the nucleic acid molecules of the present invention include nucleic acid molecules that encode a protein or fragment thereof. Another subset of the nucleic acid molecules of the present invention are EST molecules.

Fragment nucleic acid molecules may encode significant portion(s) of, or indeed most of, these nucleic acid molecules. Alternatively, the fragments may comprise smaller oligonucleotides (having from about 15 to about 250 nucleotide residues and more preferably, about 15 to about 30 nucleotide residues).

As used herein, an agent, be it a naturally occurring molecule or otherwise may be “substantially purified,” if desired, such that one or more molecules that is or may be present in a naturally occurring preparation containing that molecule will have been removed or will be present at a lower concentration than that at which it would normally be found.

The agents of the present invention will preferably be “biologically active” with respect to either a structural attribute, such as the capacity of a nucleic acid to hybridize to another nucleic acid molecule, or the ability of a protein to be bound by an antibody (or to compete with another molecule for such binding). Alternatively, such an attribute may be catalytic and thus involve the capacity of the agent to mediate a chemical reaction or response.

The agents of the present invention may also be recombinant. As used herein, the term recombinant means any agent (e.g. DNA, peptide etc.), that is, or results, however indirect, from human manipulation of a nucleic acid molecule.

It is understood that the agents of the present invention may be labeled with reagents that facilitate detection of the agent (e.g. fluorescent labels, Prober *et al.*, *Science* 238:336-340 (1987); Albarella *et al.*, EP 144914; chemical labels, Sheldon *et al.*, U.S. Patent 4,582,789; Albarella *et al.*, U.S. Patent 4,563,417; modified bases, Miyoshi *et al.*, EP 119448, all of which are hereby incorporated by reference in their entirety).

It is further understood, that the present invention provides recombinant bacterial, mammalian, microbial, insect, fungal and plant cells and viral constructs comprising the agents of the present invention. (See, for example, Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants; Section (b) Fungal Constructs and Fungal Transformants; Section (c) Mammalian Constructs and Transformed Mammalian Cells; Section (d) Insect Constructs and Transformed Insect Cells; and Section (e) Bacterial Constructs and Transformed Bacterial Cells)

Nucleic acid molecules or fragments thereof of the present invention are capable of specifically hybridizing to other nucleic acid molecules under certain circumstances. As used herein, two nucleic acid molecules are said to be capable of specifically hybridizing to one another if the two molecules are capable of forming an anti-parallel, double-stranded nucleic acid structure. A nucleic acid molecule is said to be the “complement” of another nucleic acid molecule if they exhibit complete complementarity. As used herein, molecules are said to exhibit “complete complementarity” when every nucleotide of one of the molecules is complementary to a nucleotide of the other. Two molecules are said to be “minimally

complementary” if they can hybridize to one another with sufficient stability to permit them to remain annealed to one another under at least conventional "low-stringency" conditions.

Similarly, the molecules are said to be “complementary” if they can hybridize to one another with sufficient stability to permit them to remain annealed to one another under conventional "high-stringency" conditions. Conventional stringency conditions are described by Sambrook *et al.*, *Molecular Cloning*, A Laboratory Manual, 2nd Ed., Cold Spring Harbor Press, Cold Spring Harbor, New York (1989) and by Haymes *et al.*, *Nucleic Acid Hybridization, A Practical Approach*, IRL Press, Washington, DC (1985), the entirety of which is herein incorporated by reference. Departures from complete complementarity are therefore permissible, as long as such departures do not completely preclude the capacity of the molecules to form a double-stranded structure. Thus, in order for a nucleic acid molecule to serve as a primer or probe it need only be sufficiently complementary in sequence to be able to form a stable double-stranded structure under the particular solvent and salt concentrations employed.

Appropriate stringency conditions which promote DNA hybridization, for example, 6.0 X sodium chloride/sodium citrate (SSC) at about 45°C, followed by a wash of 2.0 X SSC at 50°C, are known to those skilled in the art or can be found in *Current Protocols in Molecular Biology*, John Wiley & Sons, N.Y. (1989), 6.3.1-6.3.6. For example, the salt concentration in the wash step can be selected from a low stringency of about 2.0 X SSC at 50°C to a high stringency of about 0.2 X SSC at 50°C. In addition, the temperature in the wash step can be increased from low stringency conditions at room temperature, about 22°C, to high stringency conditions at about 65°C. Both temperature and salt may be varied, or either the temperature or the salt concentration may be held constant while the other variable is changed.

In a preferred embodiment, a nucleic acid of the present invention will specifically hybridize to one or more of the nucleic acid molecules set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof under moderately stringent conditions, for example at about 2.0 X SSC and about 65°C.

In a particularly preferred embodiment, a nucleic acid of the present invention will include those nucleic acid molecules that specifically hybridize to one or more of the nucleic acid molecules set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof under high stringency conditions such as 0.2 X SSC and about 65°C.

In one aspect of the present invention, the nucleic acid molecules of the present invention have one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In another aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 90% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In a further aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 95% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In a more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 98% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In an even more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 99% sequence identity with one or more of the sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof.

In a further more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention exhibit 100% sequence identity with a nucleic acid molecule present within MONN01, SATMON001 through SATMON031, SATMON033, SATMON034, SATMON~001, SATMONN01, SATMONN04 through SATMONN006, CMz029 through CMz031, CMz033, CMz035 through CMz037, CMz039 through CMz042, CMz044 through CMz045, CMz047 through CMz050, SOYMON001 through SOYMON038, Soy51 through Soy56, Soy58 through Soy62, Soy65 through Soy66, Soy 68 through Soy73 and Soy76 through Soy77, Lib9, Lib22 through Lib25, Lib35, Lib80 through Lib81, Lib 144, Lib146, Lib147, Lib190, Lib3032 through Lib3036 and Lib3099 (Monsanto Company, St. Louis, Missouri U.S.A.).

(i) Nucleic Acid Molecules Encoding Proteins or Fragments Thereof

Nucleic acid molecules of the present invention can comprise sequences that encode a methionine pathway protein or fragment thereof. Such proteins or fragments thereof include homologues of known proteins in other organisms.

In a preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of another plant protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of a fungal protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue of the present invention is a homologue of mammalian protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of a bacterial protein. In another preferred embodiment of the present invention, a soybean protein homologue or fragment thereof of the present invention is a

maize protein homologue or fragment thereof of the present invention is a homologue of a soybean protein.

In a preferred embodiment of the present invention, the nucleic molecule of the present invention encodes a maize or soybean homologue protein or fragment thereof where a maize or soybean homologue protein exhibits a BLAST probability score of greater than 1E-12, preferably a BLAST probability score of between about 1E-30 and about 1E-12, even more preferably a BLAST probability score of greater than 1E-30 with its homologue.

In another preferred embodiment of the present invention, the nucleic acid molecule encoding a maize or soybean protein homologue or fragment thereof or fragment thereof exhibits a % identity with its homologue of between about 25% and about 40%, more preferably of between about 40 and about 70%, even more preferably of between about 70% and about 90% and even more preferably between about 90% and 99%. In another preferred embodiment, of the present invention, a maize or soybean protein homologue or fragments thereof exhibits a % identity with its homologue of 100%.

In a preferred embodiment of the present invention, the nucleic molecule of the present invention encodes a maize or soybean homologue protein or fragment thereof where a maize or soybean homologue protein exhibits a BLAST score of greater than 120, preferably a BLAST score of between about 1450 and about 120, even more preferably a BLAST score of greater than 1450 with its homologue.

Nucleic acid molecules of the present invention also include non-maize, non-soybean homologues. Preferred non- homologues are selected from the group consisting of alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry,

sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm and *Phaseolus*.

In a preferred embodiment, nucleic acid molecules having SEQ ID NO: 1 through SEQ ID NO: 3204 or complements and fragments of either can be utilized to obtain such homologues.

The degeneracy of the genetic code, which allows different nucleic acid sequences to code for the same protein or peptide, is known in the literature. (U.S. Patent No. 4,757,006, the entirety of which is herein incorporated by reference).

In an aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding maize or soybean homologue or fragment thereof in SEQ ID NO: 1 through SEQ ID NO: 3204 due to the degeneracy in the genetic code in that they encode the same protein but differ in nucleic acid sequence.

In another further aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding maize or soybean homologue or fragment thereof in SEQ ID NO: 1 through SEQ ID NO: 3204 due to fact that the different nucleic acid sequence encodes a protein having one or more conservative amino acid residue. Examples of conservative substitutions are set forth in Table 1. It is understood that codons capable of coding for such conservative substitutions are known in the art.

Table 1

<u>Original Residue</u>	<u>Conservative Substitutions</u>
Ala	Ser
Arg	Lys

Asn	Gln; His
Asp	Glu
Cys	Ser; Ala
Gln	Asn
Glu	Asp
Gly	Pro
His	Asn; Gln
Ile	Leu; Val
Leu	Ile; Val
Lys	Arg; Gln; Glu
Met	Leu; Ile
Phe	Met; Leu; Tyr
Ser	Thr
Thr	Ser
Trp	Tyr
Tyr	Trp; Phe
Val	Ile; Leu

In a further aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding a maize or soybean homologue or fragment thereof set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof due to the fact that one or more codons encoding an amino acid has been

substituted for a codon that encodes a nonessential substitution of the amino acid originally encoded.

Agents of the present invention include nucleic acid molecules that encode a maize or soybean methionine pathway protein or fragment thereof and particularly substantially purified nucleic acid molecules selected from the group consisting of a nucleic acid molecule that encodes a maize or soybean methionine adenosyltransferase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean S-adenosylmethionine decarboxylase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean aspartate kinase protein or fragment thereof, a nucleic acid molecule that encode a maize or soybean aspartate-semialdehyde dehydrogenase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine β -lyase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean adenosylhomocysteine protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine β -synthase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine γ -lyase protein or fragment thereof, and a nucleic acid molecule that encodes a maize or soybean *O*-acetylhomoserine (thiol)-lyase protein or fragment thereof.

Non-limiting examples of such nucleic acid molecules of the present invention are nucleic acid molecules comprising: SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof that encode for a methionine pathway protein or fragment thereof, SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or fragment thereof that

encode for a methionine adenosyltransferase protein or fragment thereof, SEQ ID NO: 430
 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623 or fragment thereof
 that encode for a S-adenosylmethionine decarboxylase protein or fragment thereof, SEQ ID NO:
 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or fragment
 thereof that encode for a aspartate kinase protein or fragment thereof, SEQ ID NO: 901 through
 SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or fragment thereof that
 encode for a aspartate-semialdehyde dehydrogenase protein or fragment thereof, SEQ ID NO:
 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment
 thereof that encode for a *O*-succinylhomoserine (thiol)-lyase protein or fragment thereof, SEQ ID
 NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or
 fragment thereof that encode for a cystathionine β -lyase protein or fragment thereof, SEQ ID
 NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or
 fragment thereof that encode for a 5-methyltetrahydropteroyltriglutamate-homocysteine-S-
 methyltransferase protein or fragment thereof, SEQ ID NO: 1354 through SEQ ID NO: 1630 and
 SEQ ID NO: 2993 through SEQ ID NO: 3199 or fragment thereof that encode for an
 adenosylhomocysteinase protein or fragment thereof, SEQ ID NO: 1631 through SEQ ID NO:
 1632 or fragment thereof that encode for a cystathionine β -synthase protein or fragment thereof,
 SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204
 or fragment thereof that encode for a cystathionine γ -lyase protein or fragment thereof, and SEQ
 ID NO: 3200 through SEQ ID NO: 3202 or fragment thereof that encode for an *O*-
 acetylhomoserine (thiol)-lyase protein or fragment thereof.

A nucleic acid molecule of the present invention can also encode an homologue of a maize or soybean methionine adenosyltransferase or fragment thereof, a maize or soybean S-adenosylmethionine decarboxylase or fragment thereof, a maize or soybean aspartate kinase or fragment thereof, a maize or soybean aspartate-semialdehyde dehydrogenase or fragment thereof, a maize or soybean *O*-succinylhomoserine (thiol)-lyase or fragment thereof, a maize or soybean cystathionine β -lyase or fragment thereof, a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof, a maize or soybean adenosylhomocysteinease or fragment thereof, a maize or soybean cystathionine β -synthase or fragment thereof, a maize or soybean cystathionine γ -lyase or fragment thereof or a maize or soybean *O*-acetylhomoserine (thiol)-lyase or fragment thereof. As used herein a homologue protein molecule or fragment thereof is a counterpart protein molecule or fragment thereof in a second species (*e.g.*, maize methionine adenosyltransferase protein is a homologue of *Arabidopsis*' methionine adenosyltransferase protein).

(ii) Nucleic Acid Molecule Markers and Probes

One aspect of the present invention concerns markers that include nucleic acid molecules SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either that can act as markers. Genetic markers of the present invention include “dominant” or “codominant” markers. “Codominant markers” reveal the presence of two or more alleles (two per diploid individual) at a locus. “Dominant markers” reveal the presence of only a single allele per locus. The presence of the dominant marker phenotype (*e.g.*, a band of DNA) is an indication that one allele is present in either the homozygous or heterozygous condition. The absence of the dominant marker phenotype (*e.g.* absence of a DNA band) is merely evidence that “some other”

undefined allele is present. In the case of populations where individuals are predominantly homozygous and loci are predominately dimorphic, dominant and codominant markers can be equally valuable. As populations become more heterozygous and multi-allelic, codominant markers often become more informative of the genotype than dominant markers. Marker molecules can be, for example, capable of detecting polymorphisms such as single nucleotide polymorphisms (SNPs).

SNPs are single base changes in genomic DNA sequence. They occur at greater frequency and are spaced with a greater uniformity throughout a genome than other reported forms of polymorphism. The greater frequency and uniformity of SNPs means that there is greater probability that such a polymorphism will be found near or in a genetic locus of interest than would be the case for other polymorphisms. SNPs are located in protein-coding regions and noncoding regions of a genome. Some of these SNPs may result in defective or variant protein expression (e.g., as a results of mutations or defective splicing). Analysis (genotyping) of characterized SNPs can require only a plus/minus assay rather than a lengthy measurement, permitting easier automation.

SNPs can be characterized using any of a variety of methods. Such methods include the direct or indirect sequencing of the site, the use of restriction enzymes (Botstein *et al.*, *Am. J. Hum. Genet.* 32:314-331 (1980), the entirety of which is herein incorporated reference; Konieczny and Ausubel, *Plant J.* 4:403-410 (1993), the entirety of which is herein incorporated by reference), enzymatic and chemical mismatch assays (Myers *et al.*, *Nature* 313:495-498 (1985), the entirety of which is herein incorporated by reference), allele-specific PCR (Newton *et al.*, *Nucl. Acids Res.* 17:2503-2516 (1989), the entirety of which is herein incorporated by reference; Wu *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:2757-2760 (1989), the entirety of which

is herein incorporated by reference), ligase chain reaction (Barany, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:189-193 (1991), the entirety of which is herein incorporated by reference), single-strand conformation polymorphism analysis (Labrune *et al.*, *Am. J. Hum. Genet.* 48: 1115-1120 (1991), the entirety of which is herein incorporated by reference), primer-directed nucleotide incorporation assays (Kuppuswami *et al.*, *Proc. Natl. Acad. Sci. USA* 88:1143-1147 (1991), the entirety of which is herein incorporated by reference), dideoxy fingerprinting (Sarkar *et al.*, *Genomics* 13:441-443 (1992), the entirety of which is herein incorporated by reference), solid-phase ELISA-based oligonucleotide ligation assays (Nikiforov *et al.*, *Nucl. Acids Res.* 22:4167-4175 (1994), the entirety of which is herein incorporated by reference), oligonucleotide fluorescence-quenching assays (Livak *et al.*, *PCR Methods Appl.* 4:357-362 (1995), the entirety of which is herein incorporated by reference), 5'-nuclease allele-specific hybridization TaqMan assay (Livak *et al.*, *Nature Genet.* 9:341-342 (1995), the entirety of which is herein incorporated by reference), template-directed dye-terminator incorporation (TDI) assay (Chen and Kwok, *Nucl. Acids Res.* 25:347-353 (1997), the entirety of which is herein incorporated by reference), allele-specific molecular beacon assay (Tyagi *et al.*, *Nature Biotech.* 16: 49-53 (1998), the entirety of which is herein incorporated by reference), PinPoint assay (Haff and Smirnov, *Genome Res.* 7: 378-388 (1997), the entirety of which is herein incorporated by reference) and dCAPS analysis (Neff *et al.*, *Plant J.* 14:387-392 (1998), the entirety of which is herein incorporated by reference).

Additional markers, such as AFLP markers, RFLP markers and RAPD markers, can be utilized (Walton, *Seed World* 22-29 (July, 1993), the entirety of which is herein incorporated by reference; Burow and Blake, *Molecular Dissection of Complex Traits*, 13-29, Paterson (ed.), CRC Press, New York (1988), the entirety of which is herein incorporated by reference). DNA

markers can be developed from nucleic acid molecules using restriction endonucleases, the PCR and/or DNA sequence information. RFLP markers result from single base changes or insertions/deletions. These codominant markers are highly abundant in plant genomes, have a medium level of polymorphism and are developed by a combination of restriction endonuclease digestion and Southern blotting hybridization. CAPS are similarly developed from restriction nuclease digestion but only of specific PCR products. These markers are also codominant, have a medium level of polymorphism and are highly abundant in the genome. The CAPS result from single base changes and insertions/deletions.

Another marker type, RAPDs, are developed from DNA amplification with random primers and result from single base changes and insertions/deletions in plant genomes. They are dominant markers with a medium level of polymorphisms and are highly abundant. AFLP markers require using the PCR on a subset of restriction fragments from extended adapter primers. These markers are both dominant and codominant are highly abundant in genomes and exhibit a medium level of polymorphism.

SSRs require DNA sequence information. These codominant markers result from repeat length changes, are highly polymorphic and do not exhibit as high a degree of abundance in the genome as CAPS, AFLPs and RAPDs SNPs also require DNA sequence information. These codominant markers result from single base substitutions. They are highly abundant and exhibit a medium of polymorphism (Rafalski *et al.*, In: *Nonmammalian Genomic Analysis*, Birren and Lai (ed.), Academic Press, San Diego, CA, pp. 75-134 (1996), the entirety of which is herein incorporated by reference). It is understood that a nucleic acid molecule of the present invention may be used as a marker.

A PCR probe is a nucleic acid molecule capable of initiating a polymerase activity while in a double-stranded structure to with another nucleic acid. Various methods for determining the structure of PCR probes and PCR techniques exist in the art. Computer generated searches using programs such as Primer3 (www-genome.wi.mit.edu/cgi-bin/primer/primer3.cgi), STSPipeline (www-genome.wi.mit.edu/cgi-bin/www-STS_Pipeline), or GeneUp (Pesole *et al.*, *BioTechniques* 25:112-123 (1998) the entirety of which is herein incorporated by reference), for example, can be used to identify potential PCR primers.

It is understood that a fragment of one or more of the nucleic acid molecules of the present invention may be a probe and specifically a PCR probe.

(b) Protein and Peptide Molecules

A class of agents comprises one or more of the protein or fragments thereof or peptide molecules encoded by SEQ ID NO: 1 through SEQ ID NO: 3204 or one or more of the protein or fragment thereof and peptide molecules encoded by other nucleic acid agents of the present invention. As used herein, the term “protein molecule” or “peptide molecule” includes any molecule that comprises five or more amino acids. It is well known in the art that proteins may undergo modification, including post-translational modifications, such as, but not limited to, disulfide bond formation, glycosylation, phosphorylation, or oligomerization. Thus, as used herein, the term “protein molecule” or “peptide molecule” includes any protein molecule that is modified by any biological or non-biological process. The terms "amino acid" and "amino acids" refer to all naturally occurring L-amino acids. This definition is meant to include norleucine, ornithine, homocysteine and homoserine.

Non-limiting examples of the protein or fragment thereof of the present invention include a maize or soybean methionine pathway protein or fragment thereof; a maize or soybean

methionine adenosyltransferase or fragment thereof, a maize or soybean S-adenosylmethionine decarboxylase or fragment thereof, a maize or soybean aspartate kinase or fragment thereof, a maize or soybean aspartate-semialdehyde dehydrogenase or fragment thereof, a maize or soybean *O*-succinylhomoserine (thiol)-lyase or fragment thereof, a maize or soybean cystathionine β -lyase or fragment thereof, a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof, a maize or soybean adenosylhomocysteinase or fragment thereof, a maize or soybean cystathionine β -synthase or fragment thereof, a maize or soybean cystathionine γ -lyase or fragment thereof or a maize or soybean *O*-acetylhomoserine (thiol)-lyase or fragment thereof.

Non-limiting examples of the protein or fragment molecules of the present invention are an methionine pathway protein or fragment thereof encoded by: SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof that encode for a methionine pathway protein or fragment thereof, SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or fragment thereof that encode for a methionine adenosyltransferase protein or fragment thereof, SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623 or fragment thereof that encode for a S-adenosylmethionine decarboxylase protein or fragment thereof, SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or fragment thereof that encode for a aspartate kinase protein or fragment thereof, SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or fragment thereof that encode for a aspartate-semialdehyde dehydrogenase protein or fragment thereof, SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment thereof that encode for a *O*-succinylhomoserine (thiol)-lyase protein or

fragment thereof, SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment thereof that encode for a cystathionine β -lyase or fragment thereof, SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or fragment thereof that encode for a 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase protein or fragment thereof, SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199 or fragment thereof that encode for an adenosylhomocysteinase protein or fragment thereof, SEQ ID NO: 1631 through SEQ ID NO: 1632 or fragment thereof that encode for a cystathionine β -synthase protein or fragment thereof, SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204 or fragment thereof that encode for a cystathionine γ -lyase protein or fragment thereof, and SEQ ID NO: 3200 through SEQ ID NO: 3202 or fragment thereof that encode for an *O*-acetylhomoserine (thiol)-lyase protein or fragment thereof.

One or more of the protein or fragment of peptide molecules may be produced via chemical synthesis, or more preferably, by expressing in a suitable bacterial or eucaryotic host. Suitable methods for expression are described by Sambrook *et al.*, (In: *Molecular Cloning, A Laboratory Manual, 2nd Edition, Cold Spring Harbor Press, Cold Spring Harbor, New York* (1989)), or similar texts. For example, the protein may be expressed in, for example, Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants; Section (b) Fungal Constructs and Fungal Transformants; Section (c) Mammalian Constructs and Transformed Mammalian Cells; Section (d) Insect Constructs and Transformed Insect Cells; and Section (e) Bacterial Constructs and Transformed Bacterial Cells.

A “protein fragment” is a peptide or polypeptide molecule whose amino acid sequence comprises a subset of the amino acid sequence of that protein. A protein or fragment thereof that comprises one or more additional peptide regions not derived from that protein is a “fusion” protein. Such molecules may be derivatized to contain carbohydrate or other moieties (such as keyhole limpet hemocyanin, etc.). Fusion protein or peptide molecules of the present invention are preferably produced via recombinant means.

Another class of agents comprise protein or peptide molecules or fragments or fusions thereof encoded by SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof in which conservative, non-essential or non-relevant amino acid residues have been added, replaced or deleted. Computerized means for designing modifications in protein structure are known in the art (Dahiyat and Mayo, *Science* 278:82-87 (1997), the entirety of which is herein incorporated by reference).

The protein molecules of the present invention include plant homologue proteins. An example of such a homologue is a homologue protein of a non-maize or non soybean plant species, that include but not limited to alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry, sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm, *Phaseolus* etc. Particularly preferred non-maize or non-soybean for use for the isolation of homologs would include, *Arabidopsis*, barley, cotton, oat, oilseed rape, rice, canola, ornamentals, sugarcane, sugarbeet, tomato, potato, wheat and turf grasses. Such a homologue can be obtained by any of a variety of methods. Most preferably, as indicated above, one or more of the disclosed sequences (SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof) will be

used to define a pair of primers that may be used to isolate the homologue-encoding nucleic acid molecules from any desired species. Such molecules can be expressed to yield homologues by recombinant means.

(c) Antibodies

One aspect of the present invention concerns antibodies, single-chain antigen binding molecules, or other proteins that specifically bind to one or more of the protein or peptide molecules of the present invention and their homologues, fusions or fragments. Such antibodies may be used to quantitatively or qualitatively detect the protein or peptide molecules of the present invention. As used herein, an antibody or peptide is said to “specifically bind” to a protein or peptide molecule of the present invention if such binding is not competitively inhibited by the presence of non-related molecules.

Nucleic acid molecules that encode all or part of the protein of the present invention can be expressed, via recombinant means, to yield protein or peptides that can in turn be used to elicit antibodies that are capable of binding the expressed protein or peptide. Such antibodies may be used in immunoassays for that protein. Such protein-encoding molecules, or their fragments may be a “fusion” molecule (i.e., a part of a larger nucleic acid molecule) such that, upon expression, a fusion protein is produced. It is understood that any of the nucleic acid molecules of the present invention may be expressed, via recombinant means, to yield proteins or peptides encoded by these nucleic acid molecules.

The antibodies that specifically bind proteins and protein fragments of the present invention may be polyclonal or monoclonal and may comprise intact immunoglobulins, or antigen binding portions of immunoglobulins fragments (such as $F(ab')$, $F(ab')_2$), or single-chain

immunoglobulins producible, for example, via recombinant means. It is understood that practitioners are familiar with the standard resource materials which describe specific conditions and procedures for the construction, manipulation and isolation of antibodies (*see*, for example, Harlow and Lane, In: *Antibodies: A Laboratory Manual*, Cold Spring Harbor Press, Cold Spring Harbor, New York (1988), the entirety of which is herein incorporated by reference).

Murine monoclonal antibodies are particularly preferred. BALB/c mice are preferred for this purpose, however, equivalent strains may also be used. The animals are preferably immunized with approximately 25 µg of purified protein (or fragment thereof) that has been emulsified in a suitable adjuvant (such as TiterMax adjuvant (Vaxcel, Norcross, GA)). Immunization is preferably conducted at two intramuscular sites, one intraperitoneal site and one subcutaneous site at the base of the tail. An additional i.v. injection of approximately 25 µg of antigen is preferably given in normal saline three weeks later. After approximately 11 days following the second injection, the mice may be bled and the blood screened for the presence of anti-protein or peptide antibodies. Preferably, a direct binding Enzyme-Linked Immunoassay (ELISA) is employed for this purpose.

More preferably, the mouse having the highest antibody titer is given a third i.v. injection of approximately 25 µg of the same protein or fragment. The splenic leukocytes from this animal may be recovered 3 days later and then permitted to fuse, most preferably, using polyethylene glycol, with cells of a suitable myeloma cell line (such as, for example, the P3X63Ag8.653 myeloma cell line). Hybridoma cells are selected by culturing the cells under “HAT” (hypoxanthine-aminopterin-thymine) selection for about one week. The resulting clones may then be screened for their capacity to produce monoclonal antibodies (“mAbs”), preferably by direct ELISA.

In one embodiment, anti-protein or peptide monoclonal antibodies are isolated using a fusion of a protein or peptide of the present invention, or conjugate of a protein or peptide of the present invention, as immunogens. Thus, for example, a group of mice can be immunized using a fusion protein emulsified in Freund's complete adjuvant (*e.g.* approximately 50 µg of antigen per immunization). At three week intervals, an identical amount of antigen is emulsified in Freund's incomplete adjuvant and used to immunize the animals. Ten days following the third immunization, serum samples are taken and evaluated for the presence of antibody. If antibody titers are too low, a fourth booster can be employed. Polysera capable of binding the protein or peptide can also be obtained using this method.

In a preferred procedure for obtaining monoclonal antibodies, the spleens of the above-described immunized mice are removed, disrupted and immune splenocytes are isolated over a ficoll gradient. The isolated splenocytes are fused, using polyethylene glycol with BALB/c-derived HGPRT (hypoxanthine guanine phosphoribosyl transferase) deficient P3x63xAg8.653 plasmacytoma cells. The fused cells are plated into 96 well microtiter plates and screened for hybridoma fusion cells by their capacity to grow in culture medium supplemented with hypoxanthine, aminopterin and thymidine for approximately 2-3 weeks.

Hybridoma cells that arise from such incubation are preferably screened for their capacity to produce an immunoglobulin that binds to a protein of interest. An indirect ELISA may be used for this purpose. In brief, the supernatants of hybridomas are incubated in microtiter wells that contain immobilized protein. After washing, the titer of bound immunoglobulin can be determined using, for example, a goat anti-mouse antibody conjugated to horseradish peroxidase. After additional washing, the amount of immobilized enzyme is determined (for example through the use of a chromogenic substrate). Such screening is performed as quickly as possible

after the identification of the hybridoma in order to ensure that a desired clone is not overgrown by non-secreting neighbor cells. Desirably, the fusion plates are screened several times since the rates of hybridoma growth vary. In a preferred sub-embodiment, a different antigenic form may be used to screen the hybridoma. Thus, for example, the splenocytes may be immunized with one immunogen, but the resulting hybridomas can be screened using a different immunogen. It is understood that any of the protein or peptide molecules of the present invention may be used to raise antibodies.

As discussed below, such antibody molecules or their fragments may be used for diagnostic purposes. Where the antibodies are intended for diagnostic purposes, it may be desirable to derivatize them, for example with a ligand group (such as biotin) or a detectable marker group (such as a fluorescent group, a radioisotope or an enzyme).

The ability to produce antibodies that bind the protein or peptide molecules of the present invention permits the identification of mimetic compounds of those molecules. A “mimetic compound” is a compound that is not that compound, or a fragment of that compound, but which nonetheless exhibits an ability to specifically bind to antibodies directed against that compound.

It is understood that any of the agents of the present invention can be substantially purified and/or be biologically active and/or recombinant.

Uses of the Agents of the Invention

Nucleic acid molecules and fragments thereof of the present invention may be employed to obtain other nucleic acid molecules from the same species (e.g., ESTs or fragment thereof from maize may be utilized to obtain other nucleic acid molecules from maize). Such nucleic acid molecules include the nucleic acid molecules that encode the complete coding sequence of a protein and promoters and flanking sequences of such molecules. In addition, such nucleic acid

molecules include nucleic acid molecules that encode for other isozymes or gene family members. Such molecules can be readily obtained by using the above-described nucleic acid molecules or fragments thereof to screen cDNA or genomic libraries obtained from maize or soybean. Methods for forming such libraries are well known in the art.

Nucleic acid molecules and fragments thereof of the present invention may also be employed to obtain nucleic acid homologues. Such homologues include the nucleic acid molecule of other plants or other organisms (*e.g.*, alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry, sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm, *Phaseolus*, etc.) including the nucleic acid molecules that encode, in whole or in part, protein homologues of other plant species or other organisms, sequences of genetic elements such as promoters and transcriptional regulatory elements. Such molecules can be readily obtained by using the above-described nucleic acid molecules or fragments thereof to screen cDNA or genomic libraries obtained from such plant species. Methods for forming such libraries are well known in the art. Such homologue molecules may differ in their nucleotide sequences from those found in one or more of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof because complete complementarity is not needed for stable hybridization. The nucleic acid molecules of the present invention therefore also include molecules that, although capable of specifically hybridizing with the nucleic acid molecules may lack “complete complementarity.”

Any of a variety of methods may be used to obtain one or more of the above-described nucleic acid molecules (Zamechik *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 83:4143-4146 (1986), the entirety of which is herein incorporated by reference; Goodchild *et al.*, *Proc. Natl. Acad. Sci.*

(U.S.A.) 85:5507-5511 (1988), the entirety of which is herein incorporated by reference; Wickstrom *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:1028-1032 (1988), the entirety of which is herein incorporated by reference; Holt *et al.*, *Molec. Cell. Biol.* 8:963-973 (1988), the entirety of which is herein incorporated by reference; Gerwartz *et al.*, *Science* 242:1303-1306 (1988), the entirety of which is herein incorporated by reference; Anfossi *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:3379-3383 (1989), the entirety of which is herein incorporated by reference; Becker *et al.*, *EMBO J.* 8:3685-3691 (1989); the entirety of which is herein incorporated by reference).

Automated nucleic acid synthesizers may be employed for this purpose. In lieu of such synthesis, the disclosed nucleic acid molecules may be used to define a pair of primers that can be used with the polymerase chain reaction (Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.* 51:263-273 (1986); Erlich *et al.*, European Patent 50,424; European Patent 84,796; European Patent 258,017; European Patent 237,362; Mullis, European Patent 201,184; Mullis *et al.*, U.S. Patent 4,683,202; Erlich, U.S. Patent 4,582,788; and Saiki *et al.*, U.S. Patent 4,683,194, all of which are herein incorporated by reference in their entirety) to amplify and obtain any desired nucleic acid molecule or fragment.

Promoter sequence(s) and other genetic elements, including but not limited to transcriptional regulatory flanking sequences, associated with one or more of the disclosed nucleic acid sequences can also be obtained using the disclosed nucleic acid sequence provided herein. In one embodiment, such sequences are obtained by incubating EST nucleic acid molecules or preferably fragments thereof with members of genomic libraries (*e.g.* maize and soybean) and recovering clones that hybridize to the EST nucleic acid molecule or fragment thereof. In a second embodiment, methods of "chromosome walking," or inverse PCR may be used to obtain such sequences (Frohman *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:8998-9002

(1988); Ohara *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:5673-5677 (1989); Pang *et al.*, *Biotechniques* 22:1046-1048 (1977); Huang *et al.*, *Methods Mol. Biol.* 69:89-96 (1997); Huang *et al.*, *Method Mol. Biol.* 67:287-294 (1997); Benkel *et al.*, *Genet. Anal.* 13:123-127 (1996); Hartl *et al.*, *Methods Mol. Biol.* 58:293-301 (1996), all of which are herein incorporated by reference in their entirety).

The nucleic acid molecules of the present invention may be used to isolate promoters of cell enhanced, cell specific, tissue enhanced, tissue specific, developmentally or environmentally regulated expression profiles. Isolation and functional analysis of the 5' flanking promoter sequences of these genes from genomic libraries, for example, using genomic screening methods and PCR techniques would result in the isolation of useful promoters and transcriptional regulatory elements. These methods are known to those of skill in the art and have been described (See, for example, Birren *et al.*, *Genome Analysis: Analyzing DNA*, 1, (1997), Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., the entirety of which is herein incorporated by reference). Promoters obtained utilizing the nucleic acid molecules of the present invention could also be modified to affect their control characteristics. Examples of such modifications would include but are not limited to enhanced sequences as reported in Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants. Such genetic elements could be used to enhance gene expression of new and existing traits for crop improvements.

In one sub-aspect, such an analysis is conducted by determining the presence and/or identity of polymorphism(s) by one or more of the nucleic acid molecules of the present invention and more preferably one or more of the EST nucleic acid molecule or fragment thereof which are associated with a phenotype, or a predisposition to that phenotype.

Any of a variety of molecules can be used to identify such polymorphism(s). In one embodiment, one or more of the EST nucleic acid molecules (or a sub-fragment thereof) may be employed as a marker nucleic acid molecule to identify such polymorphism(s). Alternatively, such polymorphisms can be detected through the use of a marker nucleic acid molecule or a marker protein that is genetically linked to (i.e., a polynucleotide that co-segregates with) such polymorphism(s).

In an alternative embodiment, such polymorphisms can be detected through the use of a marker nucleic acid molecule that is physically linked to such polymorphism(s). For this purpose, marker nucleic acid molecules comprising a nucleotide sequence of a polynucleotide located within 1mb of the polymorphism(s) and more preferably within 100kb of the polymorphism(s) and most preferably within 10kb of the polymorphism(s) can be employed.

The genomes of animals and plants naturally undergo spontaneous mutation in the course of their continuing evolution (Gusella, *Ann. Rev. Biochem.* 55:831-854 (1986)). A “polymorphism” is a variation or difference in the sequence of the gene or its flanking regions that arises in some of the members of a species. The variant sequence and the “original” sequence co-exist in the species’ population. In some instances, such co-existence is in stable or quasi-stable equilibrium.

A polymorphism is thus said to be “allelic,” in that, due to the existence of the polymorphism, some members of a species may have the original sequence (i.e., the original “allele”) whereas other members may have the variant sequence (i.e., the variant “allele”). In the simplest case, only one variant sequence may exist and the polymorphism is thus said to be di-allelic. In other cases, the species’ population may contain multiple alleles and the polymorphism is termed tri-allelic, etc. A single gene may have multiple different unrelated

polymorphisms. For example, it may have a di-allelic polymorphism at one site and a multi-allelic polymorphism at another site.

The variation that defines the polymorphism may range from a single nucleotide variation to the insertion or deletion of extended regions within a gene. In some cases, the DNA sequence variations are in regions of the genome that are characterized by short tandem repeats (STRs) that include tandem di- or tri-nucleotide repeated motifs of nucleotides. Polymorphisms characterized by such tandem repeats are referred to as "variable number tandem repeat" ("VNTR") polymorphisms. VNTRs have been used in identity analysis (Weber, U.S. Patent 5,075,217; Armour *et al.*, *FEBS Lett.* 307:113-115 (1992); Jones *et al.*, *Eur. J. Haematol.* 39:144-147 (1987); Horn *et al.*, PCT Patent Application WO91/14003; Jeffreys, European Patent Application 370,719; Jeffreys, U.S. Patent 5,175,082; Jeffreys *et al.*, *Amer. J. Hum. Genet.* 39:11-24 (1986); Jeffreys *et al.*, *Nature* 316:76-79 (1985); Gray *et al.*, *Proc. R. Acad. Soc. Lond.* 243:241-253 (1991); Moore *et al.*, *Genomics* 10:654-660 (1991); Jeffreys *et al.*, *Anim. Genet.* 18:1-15 (1987); Hillel *et al.*, *Anim. Genet.* 20:145-155 (1989); Hillel *et al.*, *Genet.* 124:783-789 (1990), all of which are herein incorporated by reference in their entirety).

The detection of polymorphic sites in a sample of DNA may be facilitated through the use of nucleic acid amplification methods. Such methods specifically increase the concentration of polynucleotides that span the polymorphic site, or include that site and sequences located either distal or proximal to it. Such amplified molecules can be readily detected by gel electrophoresis or other means.

The most preferred method of achieving such amplification employs the polymerase chain reaction ("PCR") (Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.* 51:263-273 (1986); Erlich *et al.*, European Patent Appln. 50,424; European Patent Appln. 84,796; European

Patent Application 258,017; European Patent Appln. 237,362; Mullis, European Patent Appln. 201,184; Mullis *et al.*, U.S. Patent No. 4,683,202; Erlich, U.S. Patent No. 4,582,788; and Saiki *et al.*, U.S. Patent No. 4,683,194), using primer pairs that are capable of hybridizing to the proximal sequences that define a polymorphism in its double-stranded form.

In lieu of PCR, alternative methods, such as the "Ligase Chain Reaction" ("LCR") may be used (Barany, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:189-193 (1991), the entirety of which is herein incorporated by reference). LCR uses two pairs of oligonucleotide probes to exponentially amplify a specific target. The sequences of each pair of oligonucleotides is selected to permit the pair to hybridize to abutting sequences of the same strand of the target. Such hybridization forms a substrate for a template-dependent ligase. As with PCR, the resulting products thus serve as a template in subsequent cycles and an exponential amplification of the desired sequence is obtained.

LCR can be performed with oligonucleotides having the proximal and distal sequences of the same strand of a polymorphic site. In one embodiment, either oligonucleotide will be designed to include the actual polymorphic site of the polymorphism. In such an embodiment, the reaction conditions are selected such that the oligonucleotides can be ligated together only if the target molecule either contains or lacks the specific nucleotide that is complementary to the polymorphic site present on the oligonucleotide. Alternatively, the oligonucleotides may be selected such that they do not include the polymorphic site (see, Segev, PCT Application WO 90/01069, the entirety of which is herein incorporated by reference).

The "Oligonucleotide Ligation Assay" ("OLA") may alternatively be employed (Landegren *et al.*, *Science* 241:1077-1080 (1988), the entirety of which is herein incorporated by reference). The OLA protocol uses two oligonucleotides which are designed to be capable of

hybridizing to abutting sequences of a single strand of a target. OLA, like LCR, is particularly suited for the detection of point mutations. Unlike LCR, however, OLA results in "linear" rather than exponential amplification of the target sequence.

Nickerson *et al.*, have described a nucleic acid detection assay that combines attributes of PCR and OLA (Nickerson *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:8923-8927 (1990), the entirety of which is herein incorporated by reference). In this method, PCR is used to achieve the exponential amplification of target DNA, which is then detected using OLA. In addition to requiring multiple and separate, processing steps, one problem associated with such combinations is that they inherit all of the problems associated with PCR and OLA.

Schemes based on ligation of two (or more) oligonucleotides in the presence of nucleic acid having the sequence of the resulting "di-oligonucleotide", thereby amplifying the di-oligonucleotide, are also known (Wu *et al.*, *Genomics* 4:560-569 (1989), the entirety of which is herein incorporated by reference) and may be readily adapted to the purposes of the present invention.

Other known nucleic acid amplification procedures, such as allele-specific oligomers, branched DNA technology, transcription-based amplification systems, or isothermal amplification methods may also be used to amplify and analyze such polymorphisms (Malek *et al.*, U.S. Patent 5,130,238; Davey *et al.*, European Patent Application 329,822; Schuster *et al.*, U.S. Patent 5,169,766; Miller *et al.*, PCT Patent Application WO 89/06700; Kwoh *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:1173-1177 (1989); Gingeras *et al.*, PCT Patent Application WO 88/10315; Walker *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:392-396 (1992), all of which are herein incorporated by reference in their entirety).

The identification of a polymorphism can be determined in a variety of ways. By correlating the presence or absence of it in a plant with the presence or absence of a phenotype, it is possible to predict the phenotype of that plant. If a polymorphism creates or destroys a restriction endonuclease cleavage site, or if it results in the loss or insertion of DNA (e.g., a VNTR polymorphism), it will alter the size or profile of the DNA fragments that are generated by digestion with that restriction endonuclease. As such, individuals that possess a variant sequence can be distinguished from those having the original sequence by restriction fragment analysis. Polymorphisms that can be identified in this manner are termed “restriction fragment length polymorphisms” (“RFLPs”). RFLPs have been widely used in human and plant genetic analyses (Glassberg, UK Patent Application 2135774; Skolnick *et al.*, *Cytogen. Cell Genet.* 32:58-67 (1982); Botstein *et al.*, *Ann. J. Hum. Genet.* 32:314-331 (1980); Fischer *et al.*, (PCT Application WO90/13668); Uhlen, PCT Application WO90/11369).

Polymorphisms can also be identified by Single Strand Conformation Polymorphism (SSCP) analysis. SSCP is a method capable of identifying most sequence variations in a single strand of DNA, typically between 150 and 250 nucleotides in length (Elles, *Methods in Molecular Medicine: Molecular Diagnosis of Genetic Diseases*, Humana Press (1996), the entirety of which is herein incorporated by reference); Orita *et al.*, *Genomics* 5:874-879 (1989), the entirety of which is herein incorporated by reference). Under denaturing conditions a single strand of DNA will adopt a conformation that is uniquely dependent on its sequence conformation. This conformation usually will be different, even if only a single base is changed. Most conformations have been reported to alter the physical configuration or size sufficiently to be detectable by electrophoresis. A number of protocols have been described for SSCP including, but not limited to, Lee *et al.*, *Anal. Biochem.* 205:289-293 (1992), the entirety of

which is herein incorporated by reference; Suzuki *et al.*, *Anal. Biochem.* 192:82-84 (1991), the entirety of which is herein incorporated by reference; Lo *et al.*, *Nucleic Acids Research* 20:1005-1009 (1992), the entirety of which is herein incorporated by reference; Sarkar *et al.*, *Genomics* 13:441-443 (1992), the entirety of which is herein incorporated by reference. It is understood that one or more of the nucleic acids of the present invention, may be utilized as markers or probes to detect polymorphisms by SSCP analysis.

Polymorphisms may also be found using a DNA fingerprinting technique called amplified fragment length polymorphism (AFLP), which is based on the selective PCR amplification of restriction fragments from a total digest of genomic DNA to profile that DNA (Vos *et al.*, *Nucleic Acids Res.* 23:4407-4414 (1995), the entirety of which is herein incorporated by reference). This method allows for the specific co-amplification of high numbers of restriction fragments, which can be visualized by PCR without knowledge of the nucleic acid sequence.

AFLP employs basically three steps. Initially, a sample of genomic DNA is cut with restriction enzymes and oligonucleotide adapters are ligated to the restriction fragments of the DNA. The restriction fragments are then amplified using PCR by using the adapter and restriction sequence as target sites for primer annealing. The selective amplification is achieved by the use of primers that extend into the restriction fragments, amplifying only those fragments in which the primer extensions match the nucleotide flanking the restriction sites. These amplified fragments are then visualized on a denaturing polyacrylamide gel.

AFLP analysis has been performed on *Salix* (Beismann *et al.*, *Mol. Ecol.* 6:989-993 (1997), the entirety of which is herein incorporated by reference), *Acinetobacter* (Janssen *et al.*, *Int. J. Syst. Bacteriol.* 47:1179-1187 (1997), the entirety of which is herein incorporated by

reference), *Aeromonas popoffi* (Huys *et al.*, *Int. J. Syst. Bacteriol.* 47:1165-1171 (1997), the entirety of which is herein incorporated by reference), rice (McCouch *et al.*, *Plant Mol. Biol.* 35:89-99 (1997), the entirety of which is herein incorporated by reference; Nandi *et al.*, *Mol. Gen. Genet.* 255:1-8 (1997), the entirety of which is herein incorporated by reference; Cho *et al.*, *Genome* 39:373-378 (1996), the entirety of which is herein incorporated by reference), barley (*Hordeum vulgare*)(Simons *et al.*, *Genomics* 44:61-70 (1997), the entirety of which is herein incorporated by reference; Waugh *et al.*, *Mol. Gen. Genet.* 255:311-321 (1997), the entirety of which is herein incorporated by reference; Qi *et al.*, *Mol. Gen. Genet.* 254:330-336 (1997), the entirety of which is herein incorporated by reference; Becker *et al.*, *Mol. Gen. Genet.* 249:65-73 (1995), the entirety of which is herein incorporated by reference), potato (Van der Voort *et al.*, *Mol. Gen. Genet.* 255:438-447 (1997), the entirety of which is herein incorporated by reference; Meksem *et al.*, *Mol. Gen. Genet.* 249:74-81 (1995), the entirety of which is herein incorporated by reference), *Phytophthora infestans* (Van der Lee *et al.*, *Fungal Genet. Biol.* 21:278-291 (1997), the entirety of which is herein incorporated by reference), *Bacillus anthracis* (Keim *et al.*, *J. Bacteriol.* 179:818-824 (1997), the entirety of which is herein incorporated by reference), *Astragalus cremnophylax* (Travis *et al.*, *Mol. Ecol.* 5:735-745 (1996), the entirety of which is herein incorporated by reference), *Arabidopsis* (Cnops *et al.*, *Mol. Gen. Genet.* 253:32-41 (1996), the entirety of which is herein incorporated by reference), *Escherichia coli* (Lin *et al.*, *Nucleic Acids Res.* 24:3649-3650 (1996), the entirety of which is herein incorporated by reference), *Aeromonas* (Huys *et al.*, *Int. J. Syst. Bacteriol.* 46:572-580 (1996), the entirety of which is herein incorporated by reference), nematode (Folkertsma *et al.*, *Mol. Plant Microbe Interact.* 9:47-54 (1996), the entirety of which is herein incorporated by reference), tomato (Thomas *et al.*, *Plant J.* 8:785-794 (1995), the entirety of which is herein incorporated by reference) and human (Latorra

et al., *PCR Methods Appl.* 3:351-358 (1994), the entirety of which is herein incorporated by reference). AFLP analysis has also been used for fingerprinting mRNA (Money et al., *Nucleic Acids Res.* 24:2616-2617 (1996), the entirety of which is herein incorporated by reference; Bachem et al., *Plant J.* 9:745-753 (1996), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acids of the present invention, may be utilized as markers or probes to detect polymorphisms by AFLP analysis or for fingerprinting RNA.

Polymorphisms may also be found using random amplified polymorphic DNA (RAPD) (Williams et al., *Nucl. Acids Res.* 18:6531-6535 (1990), the entirety of which is herein incorporated by reference) and cleaveable amplified polymorphic sequences (CAPS) (Lyamichev et al., *Science* 260:778-783 (1993), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules of the present invention, may be utilized as markers or probes to detect polymorphisms by RAPD or CAPS analysis.

Through genetic mapping, a fine scale linkage map can be developed using DNA markers and, then, a genomic DNA library of large-sized fragments can be screened with molecular markers linked to the desired trait. Molecular markers are advantageous for agronomic traits that are otherwise difficult to tag, such as resistance to pathogens, insects and nematodes, tolerance to abiotic stress, quality parameters and quantitative traits such as high yield potential.

The essential requirements for marker-assisted selection in a plant breeding program are: (1) the marker(s) should co-segregate or be closely linked with the desired trait; (2) an efficient means of screening large populations for the molecular marker(s) should be available; and (3) the screening technique should have high reproducibility across laboratories and preferably be economical to use and be user-friendly.

The genetic linkage of marker molecules can be established by a gene mapping model such as, without limitation, the flanking marker model reported by Lander and Botstein, *Genetics* 121:185-199 (1989) and the interval mapping, based on maximum likelihood methods described by Lander and Botstein, *Genetics* 121:185-199 (1989) and implemented in the software package MAPMAKER/QTL (Lincoln and Lander, *Mapping Genes Controlling Quantitative Traits Using MAPMAKER/QTL*, Whitehead Institute for Biomedical Research, Massachusetts, (1990). Additional software includes Qgene, Version 2.23 (1996), Department of Plant Breeding and Biometry, 266 Emerson Hall, Cornell University, Ithaca, NY, the manual of which is herein incorporated by reference in its entirety). Use of Qgene software is a particularly preferred approach.

A maximum likelihood estimate (MLE) for the presence of a marker is calculated, together with an MLE assuming no QTL effect, to avoid false positives. A \log_{10} of an odds ratio (LOD) is then calculated as: $LOD = \log_{10} (MLE \text{ for the presence of a QTL} / MLE \text{ given no linked QTL})$.

The LOD score essentially indicates how much more likely the data are to have arisen assuming the presence of a QTL than in its absence. The LOD threshold value for avoiding a false positive with a given confidence, say 95%, depends on the number of markers and the length of the genome. Graphs indicating LOD thresholds are set forth in Lander and Botstein, *Genetics* 121:185-199 (1989) the entirety of which is herein incorporated by reference and further described by Arús and Moreno-González, *Plant Breeding*, Hayward *et al.*, (eds.) Chapman & Hall, London, pp. 314-331 (1993), the entirety of which is herein incorporated by reference.

Additional models can be used. Many modifications and alternative approaches to interval mapping have been reported, including the use non-parametric methods (Kruglyak and Lander, *Genetics* 139:1421-1428 (1995), the entirety of which is herein incorporated by reference). Multiple regression methods or models can be also be used, in which the trait is regressed on a large number of markers (Jansen, *Biometrics in Plant Breeding*, van Oijen and Jansen (eds.), Proceedings of the Ninth Meeting of the Eucarpia Section Biometrics in Plant Breeding, The Netherlands, pp. 116-124 (1994); Weber and Wricke, *Advances in Plant Breeding*, Blackwell, Berlin, 16 (1994), both of which is herein incorporated by reference in their entirety). Procedures combining interval mapping with regression analysis, whereby the phenotype is regressed onto a single putative QTL at a given marker interval and at the same time onto a number of markers that serve as 'cofactors,' have been reported by Jansen and Stam, *Genetics* 136:1447-1455 (1994), the entirety of which is herein incorporated by reference and Zeng, *Genetics* 136:1457-1468 (1994) the entirety of which is herein incorporated by reference. Generally, the use of cofactors reduces the bias and sampling error of the estimated QTL positions (Utz and Melchinger, *Biometrics in Plant Breeding*, van Oijen and Jansen (eds.) Proceedings of the Ninth Meeting of the Eucarpia Section Biometrics in Plant Breeding, The Netherlands, pp.195-204 (1994), the entirety of which is herein incorporated by reference, thereby improving the precision and efficiency of QTL mapping (Zeng, *Genetics* 136:1457-1468 (1994)). These models can be extended to multi-environment experiments to analyze genotype-environment interactions (Jansen *et al.*, *Theo. Appl. Genet.* 91:33-37 (1995), the entirety of which is herein incorporated by reference).

Selection of an appropriate mapping populations is important to map construction. The choice of appropriate mapping population depends on the type of marker systems employed

(Tanksley *et al.*, *Molecular mapping plant chromosomes. Chromosome structure and function: Impact of new concepts*, Gustafson and Appels (eds.), Plenum Press, New York, pp 157-173 (1988), the entirety of which is herein incorporated by reference). Consideration must be given to the source of parents (adapted vs. exotic) used in the mapping population. Chromosome pairing and recombination rates can be severely disturbed (suppressed) in wide crosses (adapted x exotic) and generally yield greatly reduced linkage distances. Wide crosses will usually provide segregating populations with a relatively large array of polymorphisms when compared to progeny in a narrow cross (adapted x adapted).

An F_2 population is the first generation of selfing after the hybrid seed is produced. Usually a single F_1 plant is selfed to generate a population segregating for all the genes in Mendelian (1:2:1) fashion. Maximum genetic information is obtained from a completely classified F_2 population using a codominant marker system (Mather, *Measurement of Linkage in Heredity*, Methuen and Co., (1938), the entirety of which is herein incorporated by reference). In the case of dominant markers, progeny tests (e.g. F_3 , BCF_2) are required to identify the heterozygotes, thus making it equivalent to a completely classified F_2 population. However, this procedure is often prohibitive because of the cost and time involved in progeny testing. Progeny testing of F_2 individuals is often used in map construction where phenotypes do not consistently reflect genotype (e.g. disease resistance) or where trait expression is controlled by a QTL. Segregation data from progeny test populations (e.g. F_3 or BCF_2) can be used in map construction. Marker-assisted selection can then be applied to cross progeny based on marker-trait map associations (F_2 , F_3), where linkage groups have not been completely disassociated by recombination events (i.e., maximum disequilibrium).

Recombinant inbred lines (RIL) (genetically related lines; usually $>F_5$, developed from continuously selfing F_2 lines towards homozygosity) can be used as a mapping population. Information obtained from dominant markers can be maximized by using RIL because all loci are homozygous or nearly so. Under conditions of tight linkage (i.e., about $<10\%$ recombination), dominant and co-dominant markers evaluated in RIL populations provide more information per individual than either marker type in backcross populations (Reiter *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:1477-1481 (1992), the entirety of which is herein incorporated by reference). However, as the distance between markers becomes larger (i.e., loci become more independent), the information in RIL populations decreases dramatically when compared to codominant markers.

Backcross populations (e.g., generated from a cross between a successful variety (recurrent parent) and another variety (donor parent) carrying a trait not present in the former) can be utilized as a mapping population. A series of backcrosses to the recurrent parent can be made to recover most of its desirable traits. Thus a population is created consisting of individuals nearly like the recurrent parent but each individual carries varying amounts or mosaic of genomic regions from the donor parent. Backcross populations can be useful for mapping dominant markers if all loci in the recurrent parent are homozygous and the donor and recurrent parent have contrasting polymorphic marker alleles (Reiter *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:1477-1481 (1992)). Information obtained from backcross populations using either codominant or dominant markers is less than that obtained from F_2 populations because one, rather than two, recombinant gametes are sampled per plant. Backcross populations, however, are more informative (at low marker saturation) when compared to RILs as the distance between linked loci increases in RIL populations (i.e. about 15% recombination). Increased

recombination can be beneficial for resolution of tight linkages, but may be undesirable in the construction of maps with low marker saturation.

Near-isogenic lines (NIL) created by many backcrosses to produce an array of individuals that are nearly identical in genetic composition except for the trait or genomic region under interrogation can be used as a mapping population. In mapping with NILs, only a portion of the polymorphic loci are expected to map to a selected region.

Bulk segregant analysis (BSA) is a method developed for the rapid identification of linkage between markers and traits of interest (Michelmore *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:9828-9832 (1991), the entirety of which is herein incorporated by reference). In BSA, two bulked DNA samples are drawn from a segregating population originating from a single cross. These bulks contain individuals that are identical for a particular trait (resistant or susceptible to particular disease) or genomic region but arbitrary at unlinked regions (i.e. heterozygous). Regions unlinked to the target region will not differ between the bulked samples of many individuals in BSA.

It is understood that one or more of the nucleic acid molecules of the present invention may be used as molecular markers. It is also understood that one or more of the protein molecules of the present invention may be used as molecular markers.

In accordance with this aspect of the present invention, a sample nucleic acid is obtained from plants cells or tissues. Any source of nucleic acid may be used. Preferably, the nucleic acid is genomic DNA. The nucleic acid is subjected to restriction endonuclease digestion. For example, one or more nucleic acid molecule or fragment thereof of the present invention can be used as a probe in accordance with the above-described polymorphic methods. The polymorphism obtained in this approach can then be cloned to identify the mutation at the coding

region which alters the protein's structure or regulatory region of the gene which affects its expression level.

In an aspect of the present invention, one or more of the nucleic molecules of the present invention are used to determine the level (i.e., the concentration of mRNA in a sample, etc.) in a plant (preferably maize or soybean) or pattern (i.e., the kinetics of expression, rate of decomposition, stability profile, etc.) of the expression of a protein encoded in part or whole by one or more of the nucleic acid molecule of the present invention (collectively, the "Expression Response" of a cell or tissue). As used herein, the Expression Response manifested by a cell or tissue is said to be "altered" if it differs from the Expression Response of cells or tissues of plants not exhibiting the phenotype. To determine whether a Expression Response is altered, the Expression Response manifested by the cell or tissue of the plant exhibiting the phenotype is compared with that of a similar cell or tissue sample of a plant not exhibiting the phenotype. As will be appreciated, it is not necessary to re-determine the Expression Response of the cell or tissue sample of plants not exhibiting the phenotype each time such a comparison is made; rather, the Expression Response of a particular plant may be compared with previously obtained values of normal plants. As used herein, the phenotype of the organism is any of one or more characteristics of an organism (e.g. disease resistance, pest tolerance, environmental tolerance such as tolerance to abiotic stress, male sterility, quality improvement or yield etc.). A change in genotype or phenotype may be transient or permanent. Also as used herein, a tissue sample is any sample that comprises more than one cell. In a preferred aspect, a tissue sample comprises cells that share a common characteristic (e.g. derived from root, seed, flower, leaf, stem or pollen etc.).

In one aspect of the present invention, an evaluation can be conducted to determine whether a particular mRNA molecule is present. One or more of the nucleic acid molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention are utilized to detect the presence or quantity of the mRNA species. Such molecules are then incubated with cell or tissue extracts of a plant under conditions sufficient to permit nucleic acid hybridization. The detection of double-stranded probe-mRNA hybrid molecules is indicative of the presence of the mRNA; the amount of such hybrid formed is proportional to the amount of mRNA. Thus, such probes may be used to ascertain the level and extent of the mRNA production in a plant's cells or tissues. Such nucleic acid hybridization may be conducted under quantitative conditions (thereby providing a numerical value of the amount of the mRNA present). Alternatively, the assay may be conducted as a qualitative assay that indicates either that the mRNA is present, or that its level exceeds a user set, predefined value.

A principle of *in situ* hybridization is that a labeled, single-stranded nucleic acid probe will hybridize to a complementary strand of cellular DNA or RNA and, under the appropriate conditions, these molecules will form a stable hybrid. When nucleic acid hybridization is combined with histological techniques, specific DNA or RNA sequences can be identified within a single cell. An advantage of *in situ* hybridization over more conventional techniques for the detection of nucleic acids is that it allows an investigator to determine the precise spatial population (Angerer *et al.*, *Dev. Biol.* 101:477-484 (1984), the entirety of which is herein incorporated by reference; Angerer *et al.*, *Dev. Biol.* 112:157-166 (1985), the entirety of which is herein incorporated by reference; Dixon *et al.*, *EMBO J.* 10:1317-1324 (1991), the entirety of which is herein incorporated by reference). *In situ* hybridization may be used to measure the

steady-state level of RNA accumulation. It is a sensitive technique and RNA sequences present in as few as 5-10 copies per cell can be detected (Hardin *et al.*, *J. Mol. Biol.* 202:417-431 (1989), the entirety of which is herein incorporated by reference). A number of protocols have been devised for *in situ* hybridization, each with tissue preparation, hybridization and washing conditions (Meyerowitz, *Plant Mol. Biol. Rep.* 5:242-250 (1987), the entirety of which is herein incorporated by reference; Cox and Goldberg, In: *Plant Molecular Biology: A Practical Approach*, Shaw (ed.), pp 1-35, IRL Press, Oxford (1988), the entirety of which is herein incorporated by reference; Raikhel *et al.*, *In situ RNA hybridization in plant tissues*, In: *Plant Molecular Biology Manual*, vol. B9:1-32, Kluwer Academic Publisher, Dordrecht, Belgium (1989), the entirety of which is herein incorporated by reference).

In situ hybridization also allows for the localization of proteins within a tissue or cell (Wilkinson, *In Situ Hybridization*, Oxford University Press, Oxford (1992), the entirety of which is herein incorporated by reference; Langdale, *In Situ Hybridization* In: *The Maize Handbook*, Freeling and Walbot (eds.), pp 165-179, Springer-Verlag, New York (1994), the entirety of which is herein incorporated by reference). It is understood that one or more of the molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention or one or more of the antibodies of the present invention may be utilized to detect the level or pattern of a methionine pathway protein or mRNA thereof by *in situ* hybridization.

Fluorescent *in situ* hybridization allows the localization of a particular DNA sequence along a chromosome which is useful, among other uses, for gene mapping, following chromosomes in hybrid lines or detecting chromosomes with translocations, transversions or deletions. *In situ* hybridization has been used to identify chromosomes in several plant species

(Griffor *et al.*, *Plant Mol. Biol.* 17:101-109 (1991), the entirety of which is herein incorporated by reference; Gustafson *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:1899-1902 (1990), herein incorporated by reference; Mukai and Gill, *Genome* 34:448-452 (1991), the entirety of which is herein incorporated by reference; Schwarzach and Heslop-Harrison, *Genome* 34:317-323 (1991); Wang *et al.*, *Jpn. J. Genet.* 66:313-316 (1991), the entirety of which is herein incorporated by reference; Parra and Windle, *Nature Genetics* 5:17-21 (1993), the entirety of which is herein incorporated by reference). It is understood that the nucleic acid molecules of the present invention may be used as probes or markers to localize sequences along a chromosome.

Another method to localize the expression of a molecule is tissue printing. Tissue printing provides a way to screen, at the same time on the same membrane many tissue sections from different plants or different developmental stages. Tissue-printing procedures utilize films designed to immobilize proteins and nucleic acids. In essence, a freshly cut section of a tissue is pressed gently onto nitrocellulose paper, nylon membrane or polyvinylidene difluoride membrane. Such membranes are commercially available (*e.g.* Millipore, Bedford, Massachusetts U.S.A.). The contents of the cut cell transfer onto the membrane and the contents and are immobilized to the membrane. The immobilized contents form a latent print that can be visualized with appropriate probes. When a plant tissue print is made on nitrocellulose paper, the cell walls leave a physical print that makes the anatomy visible without further treatment (Varner and Taylor, *Plant Physiol.* 91:31-33 (1989), the entirety of which is herein incorporated by reference).

Tissue printing on substrate films is described by Daoust, *Exp. Cell Res.* 12:203-211 (1957), the entirety of which is herein incorporated by reference, who detected amylase, protease, ribonuclease and deoxyribonuclease in animal tissues using starch, gelatin and agar films. These

techniques can be applied to plant tissues (Yomo and Taylor, *Planta* 112:35-43 (1973); the entirety of which is herein incorporated by reference; Harris and Chrispeels, *Plant Physiol.* 56:292-299 (1975), the entirety of which is herein incorporated by reference). Advances in membrane technology have increased the range of applications of Daoust's tissue-printing techniques allowing (Cassab and Varner, *J. Cell. Biol.* 105:2581-2588 (1987), the entirety of which is herein incorporated by reference) the histochemical localization of various plant enzymes and deoxyribonuclease on nitrocellulose paper and nylon (Spruce *et al.*, *Phytochemistry* 26:2901-2903 (1987), the entirety of which is herein incorporated by reference; Barres *et al.*, *Neuron* 5:527-544 (1990), the entirety of which is herein incorporated by reference; Reid and Pont-Lezica, *Tissue Printing: Tools for the Study of Anatomy, Histochemistry and Gene Expression*, Academic Press, New York, New York (1992), the entirety of which is herein incorporated by reference; Reid *et al.*, *Plant Physiol.* 93:160-165 (1990), the entirety of which is herein incorporated by reference; Ye *et al.*, *Plant J.* 1:175-183 (1991), the entirety of which is herein incorporated by reference).

It is understood that one or more of the molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention or one or more of the antibodies of the present invention may be utilized to detect the presence or quantity of a methionine pathway protein by tissue printing.

Further it is also understood that any of the nucleic acid molecules of the present invention may be used as marker nucleic acids and or probes in connection with methods that require probes or marker nucleic acids. As used herein, a probe is an agent that is utilized to determine an attribute or feature (e.g. presence or absence, location, correlation, etc.) of a molecule, cell, tissue or plant. As used herein, a marker nucleic acid is a nucleic acid molecule

that is utilized to determine an attribute or feature (*e.g.*, presence or absence, location, correlation, etc.) or a molecule, cell, tissue or plant.

A microarray-based method for high-throughput monitoring of plant gene expression may be utilized to measure gene-specific hybridization targets. This 'chip'-based approach involves using microarrays of nucleic acid molecules as gene-specific hybridization targets to quantitatively measure expression of the corresponding plant genes (Schena *et al.*, *Science* 270:467-470 (1995), the entirety of which is herein incorporated by reference; Shalon, Ph.D. Thesis, Stanford University (1996), the entirety of which is herein incorporated by reference). Every nucleotide in a large sequence can be queried at the same time. Hybridization can be used to efficiently analyze nucleotide sequences.

Several microarray methods have been described. One method compares the sequences to be analyzed by hybridization to a set of oligonucleotides representing all possible subsequences (Bains and Smith, *J. Theor. Biol.* 135:303-307 (1989), the entirety of which is herein incorporated by reference). A second method hybridizes the sample to an array of oligonucleotide or cDNA molecules. An array consisting of oligonucleotides complementary to subsequences of a target sequence can be used to determine the identity of a target sequence, measure its amount and detect differences between the target and a reference sequence. Nucleic acid molecules microarrays may also be screened with protein molecules or fragments thereof to determine nucleic acid molecules that specifically bind protein molecules or fragments thereof.

The microarray approach may be used with polypeptide targets (U.S. Patent No. 5,445,934; U.S. Patent No. 5,143,854; U.S. Patent No. 5,079,600; U.S. Patent No. 4,923,901, all of which are herein incorporated by reference in their entirety). Essentially, polypeptides are synthesized on a substrate (microarray) and these polypeptides can be screened with either

protein molecules or fragments thereof or nucleic acid molecules in order to screen for either protein molecules or fragments thereof or nucleic acid molecules that specifically bind the target polypeptides. (Fodor *et al.*, *Science* 251:767-773 (1991), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules or protein or fragments thereof of the present invention may be utilized in a microarray based method.

In a preferred embodiment of the present invention microarrays may be prepared that comprise nucleic acid molecules where such nucleic acid molecules encode at least one, preferably at least two, more preferably at least three, even more preferably at least four, five six or seven methionine pathway enzymes. In a preferred embodiment the nucleic acid molecules are selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a

maize or a soybean cystathionine γ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof.

Site directed mutagenesis may be utilized to modify nucleic acid sequences, particularly as it is a technique that allows one or more of the amino acids encoded by a nucleic acid molecule to be altered (e.g. a threonine to be replaced by a methionine). Three basic methods for site directed mutagenesis are often employed. These are cassette mutagenesis (Wells *et al.*, *Gene* 34:315-323 (1985), the entirety of which is herein incorporated by reference), primer extension (Gilliam *et al.*, *Gene* 12:129-137 (1980), the entirety of which is herein incorporated by reference; Zoller and Smith, *Methods Enzymol.* 100:468-500 (1983), the entirety of which is herein incorporated by reference; Dalbadie-McFarland *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 79:6409-6413 (1982), the entirety of which is herein incorporated by reference) and methods based upon PCR (Scharf *et al.*, *Science* 233:1076-1078 (1986), the entirety of which is herein incorporated by reference; Higuchi *et al.*, *Nucleic Acids Res.* 16:7351-7367 (1988), the entirety of which is herein incorporated by reference). Site directed mutagenesis approaches are also described in European Patent 0 385 962, the entirety of which is herein incorporated by reference; European Patent 0 359 472, the entirety of which is herein incorporated by reference; and PCT Patent Application WO 93/07278, the entirety of which is herein incorporated by reference.

Site directed mutagenesis strategies have been applied to plants for both *in vitro* as well as *in vivo* site directed mutagenesis (Lanz *et al.*, *J. Biol. Chem.* 266:9971-9976 (1991), the entirety of which is herein incorporated by reference; Kovgan and Zhdanov, *Biotekhnologiya* 5:148-154, No. 207160n, Chemical Abstracts 110:225 (1989), the entirety of which is herein incorporated by reference; Ge *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:4037-4041 (1989), the

entirety of which is herein incorporated by reference; Zhu *et al.*, *J. Biol. Chem.* 271:18494-18498 (1996), the entirety of which is herein incorporated by reference; Chu *et al.*, *Biochemistry* 33:6150-6157 (1994), the entirety of which is herein incorporated by reference; Small *et al.*, *EMBO J.* 11:1291-1296 (1992), the entirety of which is herein incorporated by reference; Cho *et al.*, *Mol. Biotechnol.* 8:13-16 (1997), the entirety of which is herein incorporated by reference; Kita *et al.*, *J. Biol. Chem.* 271:26529-26535 (1996), the entirety of which is herein incorporated by reference; Jin *et al.*, *Mol. Microbiol.* 7:555-562 (1993), the entirety of which is herein incorporated by reference; Hatfield and Vierstra, *J. Biol. Chem.* 267:14799-14803 (1992), the entirety of which is herein incorporated by reference; Zhao *et al.*, *Biochemistry* 31:5093-5099 (1992), the entirety of which is herein incorporated by reference).

Any of the nucleic acid molecules of the present invention may either be modified by site directed mutagenesis or used as, for example, nucleic acid molecules that are used to target other nucleic acid molecules for modification. It is understood that mutants with more than one altered nucleotide can be constructed using techniques that practitioners are familiar with such as isolating restriction fragments and ligating such fragments into an expression vector (*see, for example, Sambrook et al., Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Press (1989)).

Sequence-specific DNA-binding proteins play a role in the regulation of transcription. The isolation of recombinant cDNAs encoding these proteins facilitates the biochemical analysis of their structural and functional properties. Genes encoding such DNA-binding proteins have been isolated using classical genetics (Vollbrecht *et al.*, *Nature* 350: 241-243 (1991), the entirety of which is herein incorporated by reference) and molecular biochemical approaches, including the screening of recombinant cDNA libraries with antibodies (Landschulz *et al.*, *Genes Dev.*

2:786-800 (1988), the entirety of which is herein incorporated by reference) or DNA probes (Bodner *et al.*, *Cell* 55:505-518 (1988), the entirety of which is herein incorporated by reference). In addition, an *in situ* screening procedure has been used and has facilitated the isolation of sequence-specific DNA-binding proteins from various plant species (Gilmartin *et al.*, *Plant Cell* 4:839-849 (1992), the entirety of which is herein incorporated by reference; Schindler *et al.*, *EMBO J.* 11:1261-1273 (1992), the entirety of which is herein incorporated by reference). An *in situ* screening protocol does not require the purification of the protein of interest (Vinson *et al.*, *Genes Dev.* 2:801-806 (1988), the entirety of which is herein incorporated by reference; Singh *et al.*, *Cell* 52:415-423 (1988), the entirety of which is herein incorporated by reference).

Two steps may be employed to characterize DNA-protein interactions. The first is to identify promoter fragments that interact with DNA-binding proteins, to titrate binding activity, to determine the specificity of binding and to determine whether a given DNA-binding activity can interact with related DNA sequences (Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, 2nd edition, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1989)). Electrophoretic mobility-shift assay is a widely used assay. The assay provides a rapid and sensitive method for detecting DNA-binding proteins based on the observation that the mobility of a DNA fragment through a nondenaturing, low-ionic strength polyacrylamide gel is retarded upon association with a DNA-binding protein (Fried and Crother, *Nucleic Acids Res.* 9:6505-6525 (1981), the entirety of which is herein incorporated by reference). When one or more specific binding activities have been identified, the exact sequence of the DNA bound by the protein may be determined. Several procedures for characterizing protein/DNA-binding sites are used, including methylation and ethylation interference assays (Maxam and Gilbert, *Methods*

Enzymol. 65:499-560 (1980), the entirety of which is herein incorporated by reference; Wissman and Hillen, *Methods Enzymol.* 208:365-379 (1991), the entirety of which is herein incorporated by reference), footprinting techniques employing DNase I (Galas and Schmitz, *Nucleic Acids Res.* 5:3157-3170 (1978), the entirety of which is herein incorporated by reference), 1,10-phenanthroline-copper ion methods (Sigman *et al.*, *Methods Enzymol.* 208:414-433 (1991), the entirety of which is herein incorporated by reference) and hydroxyl radicals methods (Dixon *et al.*, *Methods Enzymol.* 208:414-433 (1991), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules of the present invention may be utilized to identify a protein or fragment thereof that specifically binds to a nucleic acid molecule of the present invention. It is also understood that one or more of the protein molecules or fragments thereof of the present invention may be utilized to identify a nucleic acid molecule that specifically binds to it.

A two-hybrid system is based on the fact that many cellular functions are carried out by proteins, such as transcription factors, that interact (physically) with one another. Two-hybrid systems have been used to probe the function of new proteins (Chien *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:9578-9582 (1991) the entirety of which is herein incorporated by reference; Durfee *et al.*, *Genes Dev.* 7:555-569 (1993) the entirety of which is herein incorporated by reference; Choi *et al.*, *Cell* 78:499-512 (1994), the entirety of which is herein incorporated by reference; Kranz *et al.*, *Genes Dev.* 8:313-327 (1994), the entirety of which is herein incorporated by reference).

Interaction mating techniques have facilitated a number of two-hybrid studies of protein-protein interaction. Interaction mating has been used to examine interactions between small sets of tens of proteins (Finley and Brent, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:12098-12984 (1994), the entirety of which is herein incorporated by reference), larger sets of hundreds of proteins

(Bendixen *et al.*, *Nucl. Acids Res.* 22:1778-1779 (1994), the entirety of which is herein incorporated by reference) and to comprehensively map proteins encoded by a small genome (Bartel *et al.*, *Nature Genetics* 12:72-77 (1996), the entirety of which is herein incorporated by reference). This technique utilizes proteins fused to the DNA-binding domain and proteins fused to the activation domain. They are expressed in two different haploid yeast strains of opposite mating type and the strains are mated to determine if the two proteins interact. Mating occurs when haploid yeast strains come into contact and result in the fusion of the two haploids into a diploid yeast strain. An interaction can be determined by the activation of a two-hybrid reporter gene in the diploid strain. An advantage of this technique is that it reduces the number of yeast transformations needed to test individual interactions. It is understood that the protein-protein interactions of protein or fragments thereof of the present invention may be investigated using the two-hybrid system and that any of the nucleic acid molecules of the present invention that encode such proteins or fragments thereof may be used to transform yeast in the two-hybrid system.

(a) Plant Constructs and Plant Transformants

One or more of the nucleic acid molecules of the present invention may be used in plant transformation or transfection. Exogenous genetic material may be transferred into a plant cell and the plant cell regenerated into a whole, fertile or sterile plant. Exogenous genetic material is any genetic material, whether naturally occurring or otherwise, from any source that is capable of being inserted into any organism. Such genetic material may be transferred into either monocotyledons and dicotyledons including, but not limited to maize (pp 63-69), soybean (pp

50-60), *Arabidopsis* (p 45), phaseolus (pp 47-49), peanut (pp 49-50), alfalfa (p 60), wheat (pp 69-71), rice (pp 72-79), oat (pp 80-81), sorghum (p 83), rye (p 84), tritordeum (p 84), millet (p85), fescue (p 85), perennial ryegrass (p 86), sugarcane (p87), cranberry (p101), papaya (pp 101-102), banana (p 103), banana (p 103), muskmelon (p 104), apple (p 104), cucumber (p 105), dendrobium (p 109), gladiolus (p 110), chrysanthemum (p 110), liliacea (p 111), cotton (pp113-114), eucalyptus (p 115), sunflower (p 118), canola (p 118), turfgrass (p121), sugarbeet (p 122), coffee (p 122) and dioscorea (p 122), (Christou, In: *Particle Bombardment for Genetic Engineering of Plants*, Biotechnology Intelligence Unit. Academic Press, San Diego, California (1996), the entirety of which is herein incorporated by reference).

Transfer of a nucleic acid that encodes for a protein can result in overexpression of that protein in a transformed cell or transgenic plant. One or more of the proteins or fragments thereof encoded by nucleic acid molecules of the present invention may be overexpressed in a transformed cell or transformed plant. Particularly, any of the methionine pathway proteins or fragments thereof may be overexpressed in a transformed cell or transgenic plant. Such overexpression may be the result of transient or stable transfer of the exogenous genetic material.

Exogenous genetic material may be transferred into a plant cell and the plant cell by the use of a DNA vector or construct designed for such a purpose. Design of such a vector is generally within the skill of the art (See, *Plant Molecular Biology: A Laboratory Manual*, Clark (ed.), Springer, New York (1997), the entirety of which is herein incorporated by reference).

A construct or vector may include a plant promoter to express the protein or protein fragment of choice. A number of promoters which are active in plant cells have been described in the literature. These include the nopaline synthase (NOS) promoter (Ebert *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 84:5745-5749 (1987), the entirety of which is herein incorporated by

reference), the octopine synthase (OCS) promoter (which are carried on tumor-inducing plasmids of *Agrobacterium tumefaciens*), the caulimovirus promoters such as the cauliflower mosaic virus (CaMV) 19S promoter (Lawton *et al.*, *Plant Mol. Biol.* 9:315-324 (1987), the entirety of which is herein incorporated by reference) and the CAMV 35S promoter (Odell *et al.*, *Nature* 313:810-812 (1985), the entirety of which is herein incorporated by reference), the figwort mosaic virus 35S-promoter, the light-inducible promoter from the small subunit of ribulose-1,5-bis-phosphate carboxylase (ssRUBISCO), the Adh promoter (Walker *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 84:6624-6628 (1987), the entirety of which is herein incorporated by reference), the sucrose synthase promoter (Yang *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:4144-4148 (1990), the entirety of which is herein incorporated by reference), the R gene complex promoter (Chandler *et al.*, *The Plant Cell* 1:1175-1183 (1989), the entirety of which is herein incorporated by reference) and the chlorophyll a/b binding protein gene promoter, etc. These promoters have been used to create DNA constructs which have been expressed in plants; *see, e.g.*, PCT publication WO 84/02913, herein incorporated by reference in its entirety.

Promoters which are known or are found to cause transcription of DNA in plant cells can be used in the present invention. Such promoters may be obtained from a variety of sources such as plants and plant viruses. It is preferred that the particular promoter selected should be capable of causing sufficient expression to result in the production of an effective amount of the methionine pathway protein to cause the desired phenotype. In addition to promoters that are known to cause transcription of DNA in plant cells, other promoters may be identified for use in the current invention by screening a plant cDNA library for genes which are selectively or preferably expressed in the target tissues or cells.

For the purpose of expression in source tissues of the plant, such as the leaf, seed, root or stem, it is preferred that the promoters utilized in the present invention have relatively high expression in these specific tissues. For this purpose, one may choose from a number of promoters for genes with tissue- or cell-specific or -enhanced expression. Examples of such promoters reported in the literature include the chloroplast glutamine synthetase GS2 promoter from pea (Edwards *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:3459-3463 (1990), herein incorporated by reference in its entirety), the chloroplast fructose-1,6-biphosphatase (FBPase) promoter from wheat (Lloyd *et al.*, *Mol. Gen. Genet.* 225:209-216 (1991), herein incorporated by reference in its entirety), the nuclear photosynthetic ST-LS1 promoter from potato (Stockhaus *et al.*, *EMBO J.* 8:2445-2451 (1989), herein incorporated by reference in its entirety), the serine/threonine kinase (PAL) promoter and the glucoamylase (CHS) promoter from *Arabidopsis thaliana*. Also reported to be active in photosynthetically active tissues are the ribulose-1,5-bisphosphate carboxylase (RbcS) promoter from eastern larch (*Larix laricina*), the promoter for the *cab* gene, *cab6*, from pine (Yamamoto *et al.*, *Plant Cell Physiol.* 35:773-778 (1994), herein incorporated by reference in its entirety), the promoter for the *Cab-1* gene from wheat (Fejes *et al.*, *Plant Mol. Biol.* 15:921-932 (1990), herein incorporated by reference in its entirety), the promoter for the *CAB-1* gene from spinach (Lubberstedt *et al.*, *Plant Physiol.* 104:997-1006 (1994), herein incorporated by reference in its entirety), the promoter for the *cab1R* gene from rice (Luan *et al.*, *Plant Cell.* 4:971-981 (1992), the entirety of which is herein incorporated by reference), the pyruvate, orthophosphate dikinase (PPDK) promoter from maize (Matsuoka *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 90: 9586-9590 (1993), herein incorporated by reference in its entirety), the promoter for the tobacco *Lhcb1*2* gene (Cerdan *et al.*, *Plant Mol. Biol.* 33:245-255 (1997), herein incorporated by reference in its entirety), the *Arabidopsis thaliana* SUC2 sucrose-

H⁺ symporter promoter (Truernit *et al.*, *Planta*. 196:564-570 (1995), herein incorporated by reference in its entirety) and the promoter for the thylakoid membrane proteins from spinach (psaD, psaF, psaE, PC, FNR, atpC, atpD, cab, rbcS). Other promoters for the chlorophyll a/b-binding proteins may also be utilized in the present invention, such as the promoters for LhcB gene and PsbP gene from white mustard (*Sinapis alba*; Kretsch *et al.*, *Plant Mol. Biol.* 28:219-229 (1995), the entirety of which is herein incorporated by reference).

For the purpose of expression in sink tissues of the plant, such as the tuber of the potato plant, the fruit of tomato, or the seed of maize, wheat, rice and barley, it is preferred that the promoters utilized in the present invention have relatively high expression in these specific tissues. A number of promoters for genes with tuber-specific or -enhanced expression are known, including the class I patatin promoter (Bevan *et al.*, *EMBO J.* 8:1899-1906 (1986); Jefferson *et al.*, *Plant Mol. Biol.* 14:995-1006 (1990), both of which are herein incorporated by reference in its entirety), the promoter for the potato tuber ADPGPP genes, both the large and small subunits, the sucrose synthase promoter (Salanoubat and Belliard, *Gene*. 60:47-56 (1987), Salanoubat and Belliard, *Gene*. 84:181-185 (1989), both of which are incorporated by reference in their entirety), the promoter for the major tuber proteins including the 22 kd protein complexes and proteinase inhibitors (Hannapel, *Plant Physiol.* 101:703-704 (1993), herein incorporated by reference in its entirety), the promoter for the granule bound starch synthase gene (GBSS) (Visser *et al.*, *Plant Mol. Biol.* 17:691-699 (1991), herein incorporated by reference in its entirety) and other class I and II patatins promoters (Koster-Topfer *et al.*, *Mol Gen Genet.* 219:390-396 (1989); Mignery *et al.*, *Gene*. 62:27-44 (1988), both of which are herein incorporated by reference in their entirety).

Other promoters can also be used to express a methionine pathway protein or fragment thereof in specific tissues, such as seeds or fruits. The promoter for β -conglycinin (Chen *et al.*, *Dev. Genet.* 10: 112-122 (1989), herein incorporated by reference in its entirety) or other seed-specific promoters such as the napin and phaseolin promoters, can be used. The zeins are a group of storage proteins found in maize endosperm. Genomic clones for zein genes have been isolated (Pedersen *et al.*, *Cell* 29:1015-1026 (1982), herein incorporated by reference in its entirety) and the promoters from these clones, including the 15 kD, 16 kD, 19 kD, 22 kD, 27 kD and γ genes, could also be used. Other promoters known to function, for example, in maize include the promoters for the following genes: *waxy*, *Brittle*, *Shrunken 2*, Branching enzymes I and II, starch synthases, debranching enzymes, oleosins, glutelins and sucrose synthases. A particularly preferred promoter for maize endosperm expression is the promoter for the glutelin gene from rice, more particularly the Osgt-1 promoter (Zheng *et al.*, *Mol. Cell Biol.* 13:5829-5842 (1993), herein incorporated by reference in its entirety). Examples of promoters suitable for expression in wheat include those promoters for the ADPGlucose pyrosynthase (ADPGPP) subunits, the granule bound and other starch synthase, the branching and debranching enzymes, the embryogenesis-abundant proteins, the gliadins and the glutenins. Examples of such promoters in rice include those promoters for the ADPGPP subunits, the granule bound and other starch synthase, the branching enzymes, the debranching enzymes, sucrose synthases and the glutelins. A particularly preferred promoter is the promoter for rice glutelin, Osgt-1. Examples of such promoters for barley include those for the ADPGPP subunits, the granule bound and other starch synthase, the branching enzymes, the debranching enzymes, sucrose synthases, the hordeins, the embryo globulins and the aleurone specific proteins.

Root specific promoters may also be used. An example of such a promoter is the promoter for the acid chitinase gene (Samac *et al.*, *Plant Mol. Biol.* 25:587-596 (1994), the entirety of which is herein incorporated by reference). Expression in root tissue could also be accomplished by utilizing the root specific subdomains of the CaMV35S promoter that have been identified (Lam *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:7890-7894 (1989), herein incorporated by reference in its entirety). Other root cell specific promoters include those reported by Conkling *et al.* (Conkling *et al.*, *Plant Physiol.* 93:1203-1211 (1990), the entirety of which is herein incorporated by reference).

Additional promoters that may be utilized are described, for example, in U.S. Patent Nos. 5,378,619; 5,391,725; 5,428,147; 5,447,858; 5,608,144; 5,608,144; 5,614,399; 5,633,441; 5,633,435; and 4,633,436, all of which are herein incorporated in their entirety. In addition, a tissue specific enhancer may be used (Fromm *et al.*, *The Plant Cell* 1:977-984 (1989), the entirety of which is herein incorporated by reference).

Constructs or vectors may also include with the coding region of interest a nucleic acid sequence that acts, in whole or in part, to terminate transcription of that region. For example, such sequences have been isolated including the Tr7 3' sequence and the NOS 3' sequence (Ingelbrecht *et al.*, *The Plant Cell* 1:671-680 (1989), the entirety of which is herein incorporated by reference; Bevan *et al.*, *Nucleic Acids Res.* 11:369-385 (1983), the entirety of which is herein incorporated by reference), or the like.

A vector or construct may also include regulatory elements. Examples of such include the Adh intron 1 (Callis *et al.*, *Genes and Develop.* 1:1183-1200 (1987), the entirety of which is herein incorporated by reference), the sucrose synthase intron (Vasil *et al.*, *Plant Physiol.* 91:1575-1579 (1989), the entirety of which is herein incorporated by reference) and the TMV

omega element (Gallie *et al.*, *The Plant Cell* 1:301-311 (1989), the entirety of which is herein incorporated by reference). These and other regulatory elements may be included when appropriate.

A vector or construct may also include a selectable marker. Selectable markers may also be used to select for plants or plant cells that contain the exogenous genetic material. Examples of such include, but are not limited to, a neo gene (Potrykus *et al.*, *Mol. Gen. Genet.* 199:183-188 (1985), the entirety of which is herein incorporated by reference) which codes for kanamycin resistance and can be selected for using kanamycin, G418, etc.; a bar gene which codes for bialaphos resistance; a mutant EPSP synthase gene (Hinchey *et al.*, *Bio/Technology* 6:915-922 (1988), the entirety of which is herein incorporated by reference) which encodes glyphosate resistance; a nitrilase gene which confers resistance to bromoxynil (Stalker *et al.*, *J. Biol. Chem.* 263:6310-6314 (1988), the entirety of which is herein incorporated by reference); a mutant acetolactate synthase gene (ALS) which confers imidazolinone or sulphonylurea resistance (European Patent Application 154,204 (Sept. 11, 1985), the entirety of which is herein incorporated by reference); and a methotrexate resistant DHFR gene (Thillet *et al.*, *J. Biol. Chem.* 263:12500-12508 (1988), the entirety of which is herein incorporated by reference).

A vector or construct may also include a transit peptide. Incorporation of a suitable chloroplast transit peptide may also be employed (European Patent Application Publication Number 0218571, the entirety of which is herein incorporated by reference). Translational enhancers may also be incorporated as part of the vector DNA. DNA constructs could contain one or more 5' non-translated leader sequences which may serve to enhance expression of the gene products from the resulting mRNA transcripts. Such sequences may be derived from the promoter selected to express the gene or can be specifically modified to increase translation of

the mRNA. Such regions may also be obtained from viral RNAs, from suitable eukaryotic genes, or from a synthetic gene sequence. For a review of optimizing expression of transgenes, see Koziel *et al.*, *Plant Mol. Biol.* 32:393-405 (1996), the entirety of which is herein incorporated by reference.

A vector or construct may also include a screenable marker. Screenable markers may be used to monitor expression. Exemplary screenable markers include a β -glucuronidase or uidA gene (GUS) which encodes an enzyme for which various chromogenic substrates are known (Jefferson, *Plant Mol. Biol. Rep.* 5:387-405 (1987), the entirety of which is herein incorporated by reference; Jefferson *et al.*, *EMBO J.* 6:3901-3907 (1987), the entirety of which is herein incorporated by reference); an R-locus gene, which encodes a product that regulates the production of anthocyanin pigments (red color) in plant tissues (Dellaporta *et al.*, *Stadler Symposium* 11:263-282 (1988), the entirety of which is herein incorporated by reference); a β -lactamase gene (Sutcliffe *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 75:3737-3741 (1978), the entirety of which is herein incorporated by reference), a gene which encodes an enzyme for which various chromogenic substrates are known (e.g., PADAC, a chromogenic cephalosporin); a luciferase gene (Ow *et al.*, *Science* 234:856-859 (1986), the entirety of which is herein incorporated by reference); a xylE gene (Zukowsky *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 80:1101-1105 (1983), the entirety of which is herein incorporated by reference) which encodes a catechol dioxygenase that can convert chromogenic catechols; an α -amylase gene (Ikata *et al.*, *Bio/Technol.* 8:241-242 (1990), the entirety of which is herein incorporated by reference); a tyrosinase gene (Katz *et al.*, *J. Gen. Microbiol.* 129:2703-2714 (1983), the entirety of which is herein incorporated by reference) which encodes an enzyme capable of oxidizing tyrosine to

DOPA and dopaquinone which in turn condenses to melanin; an α -galactosidase, which will turn a chromogenic α -galactose substrate.

Included within the terms “selectable or screenable marker genes” are also genes which encode a secretable marker whose secretion can be detected as a means of identifying or selecting for transformed cells. Examples include markers which encode a secretable antigen that can be identified by antibody interaction, or even secretable enzymes which can be detected catalytically. Secretable proteins fall into a number of classes, including small, diffusible proteins which are detectable, (*e.g.*, by ELISA), small active enzymes which are detectable in extracellular solution (*e.g.*, α -amylase, β -lactamase, phosphinothricin transferase), or proteins which are inserted or trapped in the cell wall (such as proteins which include a leader sequence such as that found in the expression unit of extension or tobacco PR-S). Other possible selectable and/or screenable marker genes will be apparent to those of skill in the art.

There are many methods for introducing transforming nucleic acid molecules into plant cells. Suitable methods are believed to include virtually any method by which nucleic acid molecules may be introduced into a cell, such as by *Agrobacterium* infection or direct delivery of nucleic acid molecules such as, for example, by PEG-mediated transformation, by electroporation or by acceleration of DNA coated particles, etc (Potrykus, *Ann. Rev. Plant Physiol. Plant Mol. Biol.* 42:205-225 (1991), the entirety of which is herein incorporated by reference; Vasil, *Plant Mol. Biol.* 25:925-937 (1994), the entirety of which is herein incorporated by reference). For example, electroporation has been used to transform maize protoplasts (Fromm *et al.*, *Nature* 312:791-793 (1986), the entirety of which is herein incorporated by reference).

Other vector systems suitable for introducing transforming DNA into a host plant cell include but are not limited to binary artificial chromosome (BIBAC) vectors (Hamilton *et al.*, *Gene* 200:107-116 (1997), the entirety of which is herein incorporated by reference); and transfection with RNA viral vectors (Della-Cioppa *et al.*, *Ann. N.Y. Acad. Sci.* (1996), 792 (Engineering Plants for Commercial Products and Applications), 57-61, the entirety of which is herein incorporated by reference). Additional vector systems also include plant selectable YAC vectors such as those described in Mullen *et al.*, *Molecular Breeding* 4:449-457 (1988), the entirety of which is herein incorporated by reference).

Technology for introduction of DNA into cells is well known to those of skill in the art. Four general methods for delivering a gene into cells have been described: (1) chemical methods (Graham and van der Eb, *Virology* 54:536-539 (1973), the entirety of which is herein incorporated by reference); (2) physical methods such as microinjection (Capecchi, *Cell* 22:479-488 (1980), the entirety of which is herein incorporated by reference), electroporation (Wong and Neumann, *Biochem. Biophys. Res. Commun.* 107:584-587 (1982); Fromm *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 82:5824-5828 (1985); U.S. Patent No. 5,384,253, all of which are herein incorporated in their entirety); and the gene gun (Johnston and Tang, *Methods Cell Biol.* 43:353-365 (1994), the entirety of which is herein incorporated by reference); (3) viral vectors (Clapp, *Clin. Perinatol.* 20:155-168 (1993); Lu *et al.*, *J. Exp. Med.* 178:2089-2096 (1993); Eglitis and Anderson, *Biotechniques* 6:608-614 (1988), all of which are herein incorporated in their entirety); and (4) receptor-mediated mechanisms (Curiel *et al.*, *Hum. Gen. Ther.* 3:147-154 (1992), Wagner *et al.*, *Proc. Natl. Acad. Sci. (USA)* 89:6099-6103 (1992), both of which are incorporated by reference in their entirety).

Acceleration methods that may be used include, for example, microprojectile bombardment and the like. One example of a method for delivering transforming nucleic acid molecules to plant cells is microprojectile bombardment. This method has been reviewed by Yang and Christou (eds.), *Particle Bombardment Technology for Gene Transfer*, Oxford Press, Oxford, England (1994), the entirety of which is herein incorporated by reference). Non-biological particles (microprojectiles) that may be coated with nucleic acids and delivered into cells by a propelling force. Exemplary particles include those comprised of tungsten, gold, platinum and the like.

A particular advantage of microprojectile bombardment, in addition to it being an effective means of reproducibly transforming monocots, is that neither the isolation of protoplasts (Cristou *et al.*, *Plant Physiol.* 87:671-674 (1988), the entirety of which is herein incorporated by reference) nor the susceptibility of *Agrobacterium* infection are required. An illustrative embodiment of a method for delivering DNA into maize cells by acceleration is a biolistics α -particle delivery system, which can be used to propel particles coated with DNA through a screen, such as a stainless steel or Nytex screen, onto a filter surface covered with corn cells cultured in suspension. Gordon-Kamm *et al.*, describes the basic procedure for coating tungsten particles with DNA (Gordon-Kamm *et al.*, *Plant Cell* 2:603-618 (1990), the entirety of which is herein incorporated by reference). The screen disperses the tungsten nucleic acid particles so that they are not delivered to the recipient cells in large aggregates. A particle delivery system suitable for use with the present invention is the helium acceleration PDS-1000/He gun is available from Bio-Rad Laboratories (Bio-Rad, Hercules, California)(Sanford *et al.*, *Technique* 3:3-16 (1991), the entirety of which is herein incorporated by reference).

For the bombardment, cells in suspension may be concentrated on filters. Filters containing the cells to be bombarded are positioned at an appropriate distance below the microprojectile stopping plate. If desired, one or more screens are also positioned between the gun and the cells to be bombarded.

Alternatively, immature embryos or other target cells may be arranged on solid culture medium. The cells to be bombarded are positioned at an appropriate distance below the microprojectile stopping plate. If desired, one or more screens are also positioned between the acceleration device and the cells to be bombarded. Through the use of techniques set forth herein one may obtain up to 1000 or more foci of cells transiently expressing a marker gene. The number of cells in a focus which express the exogenous gene product 48 hours post-bombardment often range from one to ten and average one to three.

In bombardment transformation, one may optimize the pre-bombardment culturing conditions and the bombardment parameters to yield the maximum numbers of stable transformants. Both the physical and biological parameters for bombardment are important in this technology. Physical factors are those that involve manipulating the DNA/microprojectile precipitate or those that affect the flight and velocity of either the macro- or microprojectiles. Biological factors include all steps involved in manipulation of cells before and immediately after bombardment, the osmotic adjustment of target cells to help alleviate the trauma associated with bombardment and also the nature of the transforming DNA, such as linearized DNA or intact supercoiled plasmids. It is believed that pre-bombardment manipulations are especially important for successful transformation of immature embryos.

In another alternative embodiment, plastids can be stably transformed. Methods disclosed for plastid transformation in higher plants include the particle gun delivery of DNA

containing a selectable marker and targeting of the DNA to the plastid genome through homologous recombination (Svab *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:8526-8530 (1990); Svab and Maliga, *Proc. Natl. Acad. Sci. (U.S.A.)* 90:913-917 (1993); Staub and Maliga, *EMBO J.* 12:601-606 (1993); U.S. Patents 5, 451,513 and 5,545,818, all of which are herein incorporated by reference in their entirety).

Accordingly, it is contemplated that one may wish to adjust various aspects of the bombardment parameters in small scale studies to fully optimize the conditions. One may particularly wish to adjust physical parameters such as gap distance, flight distance, tissue distance and helium pressure. One may also minimize the trauma reduction factors by modifying conditions which influence the physiological state of the recipient cells and which may therefore influence transformation and integration efficiencies. For example, the osmotic state, tissue hydration and the subculture stage or cell cycle of the recipient cells may be adjusted for optimum transformation. The execution of other routine adjustments will be known to those of skill in the art in light of the present disclosure.

Agrobacterium-mediated transfer is a widely applicable system for introducing genes into plant cells because the DNA can be introduced into whole plant tissues, thereby bypassing the need for regeneration of an intact plant from a protoplast. The use of *Agrobacterium*-mediated plant integrating vectors to introduce DNA into plant cells is well known in the art. See, for example the methods described by Fraley *et al.*, *Bio/Technology* 3:629-635 (1985) and Rogers *et al.*, *Methods Enzymol.* 153:253-277 (1987), both of which are herein incorporated by reference in their entirety. Further, the integration of the Ti-DNA is a relatively precise process resulting in few rearrangements. The region of DNA to be transferred is defined by the border sequences and

intervening DNA is usually inserted into the plant genome as described (Spielmann *et al.*, *Mol. Gen. Genet.* 205:34 (1986), the entirety of which is herein incorporated by reference).

Modern *Agrobacterium* transformation vectors are capable of replication in *E. coli* as well as *Agrobacterium*, allowing for convenient manipulations as described (Klee *et al.*, In: *Plant DNA Infectious Agents*, Hohn and Schell (eds.), Springer-Verlag, New York, pp. 179-203 (1985), the entirety of which is herein incorporated by reference. Moreover, technological advances in vectors for *Agrobacterium*-mediated gene transfer have improved the arrangement of genes and restriction sites in the vectors to facilitate construction of vectors capable of expressing various polypeptide coding genes. The vectors described have convenient multi-linker regions flanked by a promoter and a polyadenylation site for direct expression of inserted polypeptide coding genes and are suitable for present purposes (Rogers *et al.*, *Methods Enzymol.* 153:253-277 (1987)). In addition, *Agrobacterium* containing both armed and disarmed Ti genes can be used for the transformations. In those plant strains where *Agrobacterium*-mediated transformation is efficient, it is the method of choice because of the facile and defined nature of the gene transfer.

A transgenic plant formed using *Agrobacterium* transformation methods typically contains a single gene on one chromosome. Such transgenic plants can be referred to as being heterozygous for the added gene. More preferred is a transgenic plant that is homozygous for the added structural gene; *i.e.*, a transgenic plant that contains two added genes, one gene at the same locus on each chromosome of a chromosome pair. A homozygous transgenic plant can be obtained by sexually mating (selfing) an independent segregant transgenic plant that contains a single added gene, germinating some of the seed produced and analyzing the resulting plants produced for the gene of interest.

It is also to be understood that two different transgenic plants can also be mated to produce offspring that contain two independently segregating added, exogenous genes. Selfing of appropriate progeny can produce plants that are homozygous for both added, exogenous genes that encode a polypeptide of interest. Back-crossing to a parental plant and out-crossing with a non-transgenic plant are also contemplated, as is vegetative propagation.

Transformation of plant protoplasts can be achieved using methods based on calcium phosphate precipitation, polyethylene glycol treatment, electroporation and combinations of these treatments (*See, for example*, Potrykus *et al.*, *Mol. Gen. Genet.* 205:193-200 (1986); Lorz *et al.*, *Mol. Gen. Genet.* 199:178 (1985); Fromm *et al.*, *Nature* 319:791 (1986); Uchimiya *et al.*, *Mol. Gen. Genet.* 204:204 (1986); Marcotte *et al.*, *Nature* 335:454-457 (1988), all of which are herein incorporated by reference in their entirety).

Application of these systems to different plant strains depends upon the ability to regenerate that particular plant strain from protoplasts. Illustrative methods for the regeneration of cereals from protoplasts are described (Fujimura *et al.*, *Plant Tissue Culture Letters* 2:74 (1985); Toriyama *et al.*, *Theor Appl. Genet.* 205:34 (1986); Yamada *et al.*, *Plant Cell Rep.* 4:85 (1986); Abdullah *et al.*, *Biotechnolog* 4:1087 (1986), all of which are herein incorporated by reference in their entirety).

To transform plant strains that cannot be successfully regenerated from protoplasts, other ways to introduce DNA into intact cells or tissues can be utilized. For example, regeneration of cereals from immature embryos or explants can be effected as described (Vasil, *Biotechnology* 6:397 (1988), the entirety of which is herein incorporated by reference). In addition, "particle gun" or high-velocity microprojectile technology can be utilized (Vasil *et al.*, *Bio/Technology* 10:667 (1992), the entirety of which is herein incorporated by reference).

Using the latter technology, DNA is carried through the cell wall and into the cytoplasm on the surface of small metal particles as described (Klein *et al.*, *Nature* 328:70 (1987); Klein *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:8502-8505 (1988); McCabe *et al.*, *Bio/Technology* 6:923 (1988), all of which are herein incorporated by reference in their entirety). The metal particles penetrate through several layers of cells and thus allow the transformation of cells within tissue explants.

Other methods of cell transformation can also be used and include but are not limited to introduction of DNA into plants by direct DNA transfer into pollen (Zhou *et al.*, *Methods Enzymol.* 101:433 (1983); Hess *et al.*, *Intern Rev. Cytol.* 107:367 (1987); Luo *et al.*, *Plant Mol Biol. Reporter* 6:165 (1988), all of which are herein incorporated by reference in their entirety), by direct injection of DNA into reproductive organs of a plant (Pena *et al.*, *Nature* 325:274 (1987), the entirety of which is herein incorporated by reference), or by direct injection of DNA into the cells of immature embryos followed by the rehydration of desiccated embryos (Neuhaus *et al.*, *Theor. Appl. Genet.* 75:30 (1987), the entirety of which is herein incorporated by reference).

The regeneration, development and cultivation of plants from single plant protoplast transformants or from various transformed explants is well known in the art (Weissbach and Weissbach, In: *Methods for Plant Molecular Biology*, Academic Press, San Diego, CA, (1988), the entirety of which is herein incorporated by reference). This regeneration and growth process typically includes the steps of selection of transformed cells, culturing those individualized cells through the usual stages of embryonic development through the rooted plantlet stage. Transgenic embryos and seeds are similarly regenerated. The resulting transgenic rooted shoots are thereafter planted in an appropriate plant growth medium such as soil.

The development or regeneration of plants containing the foreign, exogenous gene that encodes a protein of interest is well known in the art. Preferably, the regenerated plants are self-pollinated to provide homozygous transgenic plants. Otherwise, pollen obtained from the regenerated plants is crossed to seed-grown plants of agronomically important lines. Conversely, pollen from plants of these important lines is used to pollinate regenerated plants. A transgenic plant of the present invention containing a desired polypeptide is cultivated using methods well known to one skilled in the art.

There are a variety of methods for the regeneration of plants from plant tissue. The particular method of regeneration will depend on the starting plant tissue and the particular plant species to be regenerated.

Methods for transforming dicots, primarily by use of *Agrobacterium tumefaciens* and obtaining transgenic plants have been published for cotton (U.S. Patent No. 5,004,863; U.S. Patent No. 5,159,135; U.S. Patent No. 5,518,908, all of which are herein incorporated by reference in their entirety); soybean (U.S. Patent No. 5,569,834; U.S. Patent No. 5,416,011; McCabe *et al.*, *Biotechnology* 6:923 (1988); Christou *et al.*, *Plant Physiol.* 87:671-674 (1988); all of which are herein incorporated by reference in their entirety); *Brassica* (U.S. Patent No. 5,463,174, the entirety of which is herein incorporated by reference); peanut (Cheng *et al.*, *Plant Cell Rep.* 15:653-657 (1996), McKently *et al.*, *Plant Cell Rep.* 14:699-703 (1995), all of which are herein incorporated by reference in their entirety); papaya; and pea (Grant *et al.*, *Plant Cell Rep.* 15:254-258 (1995), the entirety of which is herein incorporated by reference).

Transformation of monocotyledons using electroporation, particle bombardment and *Agrobacterium* have also been reported. Transformation and plant regeneration have been achieved in asparagus (Bytebier *et al.*, *Proc. Natl. Acad. Sci. (USA)* 84:5354 (1987), the entirety

of which is herein incorporated by reference); barley (Wan and Lemaux, *Plant Physiol* 104:37 (1994), the entirety of which is herein incorporated by reference); maize (Rhodes *et al.*, *Science* 240:204 (1988); Gordon-Kamm *et al.*, *Plant Cell* 2:603-618 (1990); Fromm *et al.*, *Bio/Technology* 8:833 (1990); Koziel *et al.*, *Bio/Technology* 11:194 (1993); Armstrong *et al.*, *Crop Science* 35:550-557 (1995); all of which are herein incorporated by reference in their entirety); oat (Somers *et al.*, *Bio/Technology* 10:1589 (1992), the entirety of which is herein incorporated by reference); orchard grass (Horn *et al.*, *Plant Cell Rep.* 7:469 (1988), the entirety of which is herein incorporated by reference); rice (Toriyama *et al.*, *Theor Appl. Genet.* 205:34 (1986); Part *et al.*, *Plant Mol. Biol.* 32:1135-1148 (1996); Abedinia *et al.*, *Aust. J. Plant Physiol.* 24:133-141 (1997); Zhang and Wu, *Theor. Appl. Genet.* 76:835 (1988); Zhang *et al.*, *Plant Cell Rep.* 7:379 (1988); Battra and Hall, *Plant Sci.* 86:191-202 (1992); Christou *et al.*, *Bio/Technology* 9:957 (1991), all of which are herein incorporated by reference in their entirety); rye (De la Pena *et al.*, *Nature* 325:274 (1987), the entirety of which is herein incorporated by reference); sugarcane (Bower and Birch, *Plant J.* 2:409 (1992), the entirety of which is herein incorporated by reference); tall fescue (Wang *et al.*, *Bio/Technology* 10:691 (1992), the entirety of which is herein incorporated by reference) and wheat (Vasil *et al.*, *Bio/Technology* 10:667 (1992), the entirety of which is herein incorporated by reference; U.S. Patent No. 5,631,152, the entirety of which is herein incorporated by reference.)

Assays for gene expression based on the transient expression of cloned nucleic acid constructs have been developed by introducing the nucleic acid molecules into plant cells by polyethylene glycol treatment, electroporation, or particle bombardment (Marcotte *et al.*, *Nature* 335:454-457 (1988), the entirety of which is herein incorporated by reference; Marcotte *et al.*, *Plant Cell* 1:523-532 (1989), the entirety of which is herein incorporated by reference; McCarty

et al., *Cell* 66:895-905 (1991), the entirety of which is herein incorporated by reference; Hattori *et al.*, *Genes Dev.* 6:609-618 (1992), the entirety of which is herein incorporated by reference; Goff *et al.*, *EMBO J.* 9:2517-2522 (1990), the entirety of which is herein incorporated by reference). Transient expression systems may be used to functionally dissect gene constructs (see generally, Mailga *et al.*, *Methods in Plant Molecular Biology*, Cold Spring Harbor Press (1995)).

Any of the nucleic acid molecules of the present invention may be introduced into a plant cell in a permanent or transient manner in combination with other genetic elements such as vectors, promoters, enhancers etc. Further, any of the nucleic acid molecules of the present invention may be introduced into a plant cell in a manner that allows for overexpression of the protein or fragment thereof encoded by the nucleic acid molecule.

Cosuppression is the reduction in expression levels, usually at the level of RNA, of a particular endogenous gene or gene family by the expression of a homologous sense construct that is capable of transcribing mRNA of the same strandedness as the transcript of the endogenous gene (Napoli *et al.*, *Plant Cell* 2:279-289 (1990), the entirety of which is herein incorporated by reference; van der Krol *et al.*, *Plant Cell* 2:291-299 (1990), the entirety of which is herein incorporated by reference). Cosuppression may result from stable transformation with a single copy nucleic acid molecule that is homologous to a nucleic acid sequence found with the cell (Prolls and Meyer, *Plant J.* 2:465-475 (1992), the entirety of which is herein incorporated by reference) or with multiple copies of a nucleic acid molecule that is homologous to a nucleic acid sequence found with the cell (Mittlesten *et al.*, *Mol. Gen. Genet.* 244:325-330 (1994), the entirety of which is herein incorporated by reference). Genes, even though different, linked to

homologous promoters may result in the cosuppression of the linked genes (Vaucheret, *C.R. Acad. Sci. III* 316:1471-1483 (1993), the entirety of which is herein incorporated by reference).

This technique has, for example, been applied to generate white flowers from red petunia and tomatoes that do not ripen on the vine. Up to 50% of petunia transformants that contained a sense copy of the glucoamylase (CHS) gene produced white flowers or floral sectors; this was as a result of the post-transcriptional loss of mRNA encoding CHS (Flavell, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:3490-3496 (1994), the entirety of which is herein incorporated by reference); van Blokland *et al.*, *Plant J.* 6:861-877 (1994), the entirety of which is herein incorporated by reference). Cosuppression may require the coordinate transcription of the transgene and the endogenous gene and can be reset by a developmental control mechanism (Jorgensen, *Trends Biotechnol.* 8:340-344 (1990), the entirety of which is herein incorporated by reference; Meins and Kunz, In: *Gene Inactivation and Homologous Recombination in Plants*, Paszkowski (ed.), pp. 335-348, Kluwer Academic, Netherlands (1994), the entirety of which is herein incorporated by reference).

It is understood that one or more of the nucleic acids of the present invention may be introduced into a plant cell and transcribed using an appropriate promoter with such transcription resulting in the cosuppression of an endogenous methionine pathway protein.

Antisense approaches are a way of preventing or reducing gene function by targeting the genetic material (Mol *et al.*, *FEBS Lett.* 268:427-430 (1990), the entirety of which is herein incorporated by reference). The objective of the antisense approach is to use a sequence complementary to the target gene to block its expression and create a mutant cell line or organism in which the level of a single chosen protein is selectively reduced or abolished. Antisense techniques have several advantages over other 'reverse genetic' approaches. The site

of inactivation and its developmental effect can be manipulated by the choice of promoter for antisense genes or by the timing of external application or microinjection. Antisense can manipulate its specificity by selecting either unique regions of the target gene or regions where it shares homology to other related genes (Hiatt *et al.*, In: *Genetic Engineering*, Setlow (ed.), Vol. 11, New York: Plenum 49-63 (1989), the entirety of which is herein incorporated by reference).

The principle of regulation by antisense RNA is that RNA that is complementary to the target mRNA is introduced into cells, resulting in specific RNA:RNA duplexes being formed by base pairing between the antisense substrate and the target mRNA (Green *et al.*, *Annu. Rev. Biochem.* 55:569-597 (1986), the entirety of which is herein incorporated by reference). Under one embodiment, the process involves the introduction and expression of an antisense gene sequence. Such a sequence is one in which part or all of the normal gene sequences are placed under a promoter in inverted orientation so that the 'wrong' or complementary strand is transcribed into a noncoding antisense RNA that hybridizes with the target mRNA and interferes with its expression (Takayama and Inouye, *Crit. Rev. Biochem. Mol. Biol.* 25:155-184 (1990), the entirety of which is herein incorporated by reference). An antisense vector is constructed by standard procedures and introduced into cells by transformation, transfection, electroporation, microinjection, infection, etc. The type of transformation and choice of vector will determine whether expression is transient or stable. The promoter used for the antisense gene may influence the level, timing, tissue, specificity, or inducibility of the antisense inhibition.

It is understood that the activity of a methionine pathway protein in a plant cell may be reduced or depressed by growing a transformed plant cell containing a nucleic acid molecule whose non-transcribed strand encodes a methionine pathway protein or fragment thereof.

Antibodies have been expressed in plants (Hiatt *et al.*, *Nature* 342:76-78 (1989), the entirety of which is herein incorporated by reference; Conrad and Fielder, *Plant Mol. Biol.* 26:1023-1030 (1994), the entirety of which is herein incorporated by reference). Cytoplasmic expression of a scFv (single-chain Fv antibodies) has been reported to delay infection by artichoke mottled crinkle virus. Transgenic plants that express antibodies directed against endogenous proteins may exhibit a physiological effect (Philips *et al.*, *EMBO J.* 16:4489-4496 (1997), the entirety of which is herein incorporated by reference; Marion-Poll, *Trends in Plant Science* 2:447-448 (1997), the entirety of which is herein incorporated by reference). For example, expressed anti-abscisic antibodies have been reported to result in a general perturbation of seed development (Philips *et al.*, *EMBO J.* 16: 4489-4496 (1997)).

Antibodies that are catalytic may also be expressed in plants (abzymes). The principle behind abzymes is that since antibodies may be raised against many molecules, this recognition ability can be directed toward generating antibodies that bind transition states to force a chemical reaction forward (Persidas, *Nature Biotechnology* 15:1313-1315 (1997), the entirety of which is herein incorporated by reference; Baca *et al.*, *Ann. Rev. Biophys. Biomol. Struct.* 26:461-493 (1997), the entirety of which is herein incorporated by reference). The catalytic abilities of abzymes may be enhanced by site directed mutagenesis. Examples of abzymes are, for example, set forth in U.S. Patent No: 5,658,753; U.S. Patent No. 5,632,990; U.S. Patent No. 5,631,137; U.S. Patent 5,602,015; U.S. Patent No. 5,559,538; U.S. Patent No. 5,576,174; U.S. Patent No. 5,500,358; U.S. Patent 5,318,897; U.S. Patent No. 5,298,409; U.S. Patent No. 5,258,289 and U.S. Patent No. 5,194,585, all of which are herein incorporated in their entirety.

It is understood that any of the antibodies of the present invention may be expressed in plants and that such expression can result in a physiological effect. It is also understood that any of the expressed antibodies may be catalytic.

(b) Fungal Constructs and Fungal Transformants

The present invention also relates to a fungal recombinant vector comprising exogenous genetic material. The present invention also relates to a fungal cell comprising a fungal recombinant vector. The present invention also relates to methods for obtaining a recombinant fungal host cell comprising introducing into a fungal host cell exogenous genetic material.

Exogenous genetic material may be transferred into a fungal cell. In a preferred embodiment the exogenous genetic material includes a nucleic acid molecule of the present invention having a sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either. The fungal recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures. The choice of a vector will typically depend on the compatibility of the vector with the fungal host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the fungal host.

The fungal vector may be an autonomously replicating vector, *i.e.*, a vector which exists as an extrachromosomal entity, the replication of which is independent of chromosomal replication, *e.g.*, a plasmid, an extrachromosomal element, a minichromosome, or an artificial chromosome. The vector may contain any means for assuring self-replication. Alternatively, the vector may be one which, when introduced into the fungal cell, is integrated into the genome and replicated together with the chromosome(s) into which it has been integrated. For integration,

the vector may rely on the nucleic acid sequence of the vector for stable integration of the vector into the genome by homologous or nonhomologous recombination. Alternatively, the vector may contain additional nucleic acid sequences for directing integration by homologous recombination into the genome of the fungal host. The additional nucleic acid sequences enable the vector to be integrated into the host cell genome at a precise location(s) in the chromosome(s). To increase the likelihood of integration at a precise location, there should be preferably two nucleic acid sequences which individually contain a sufficient number of nucleic acids, preferably 400bp to 1500bp, more preferably 800bp to 1000bp, which are highly homologous with the corresponding target sequence to enhance the probability of homologous recombination. These nucleic acid sequences may be any sequence that is homologous with a target sequence in the genome of the fungal host cell and, furthermore, may be non-encoding or encoding sequences.

For autonomous replication, the vector may further comprise an origin of replication enabling the vector to replicate autonomously in the host cell in question. Examples of origin of replications for use in a yeast host cell are the 2 micron origin of replication and the combination of CEN3 and ARS 1. Any origin of replication may be used which is compatible with the fungal host cell of choice.

The fungal vectors of the present invention preferably contain one or more selectable markers which permit easy selection of transformed cells. A selectable marker is a gene the product of which provides, for example biocide or viral resistance, resistance to heavy metals, prototrophy to auxotrophs and the like. The selectable marker may be selected from the group including, but not limited to, *amdS* (acetamidase), *argB* (ornithine carbamoyltransferase), *bar* (phosphinothricin acetyltransferase), *hygB* (hygromycin phosphotransferase), *niaD* (nitrate

reductase), *pyrG* (orotidine-5'-phosphate decarboxylase) and *sC* (sulfate adenylyltransferase) and *trpC* (anthranilate synthase). Preferred for use in an *Aspergillus* cell are the *amdS* and *pyrG* markers of *Aspergillus nidulans* or *Aspergillus oryzae* and the bar marker of *Streptomyces hygroscopicus*. Furthermore, selection may be accomplished by co-transformation, *e.g.*, as described in WO 91/17243, the entirety of which is herein incorporated by reference. A nucleic acid sequence of the present invention may be operably linked to a suitable promoter sequence. The promoter sequence is a nucleic acid sequence which is recognized by the fungal host cell for expression of the nucleic acid sequence. The promoter sequence contains transcription and translation control sequences which mediate the expression of the protein or fragment thereof.

A promoter may be any nucleic acid sequence which shows transcriptional activity in the fungal host cell of choice and may be obtained from genes encoding polypeptides either homologous or heterologous to the host cell. Examples of suitable promoters for directing the transcription of a nucleic acid construct of the invention in a filamentous fungal host are promoters obtained from the genes encoding *Aspergillus oryzae* TAKA amylase, *Rhizomucor miehei* aspartic proteinase, *Aspergillus niger* neutral alpha-amylase, *Aspergillus niger* acid stable alpha-amylase, *Aspergillus niger* or *Aspergillus awamori* glucoamylase (*glaA*), *Rhizomucor miehei* lipase, *Aspergillus oryzae* alkaline protease, *Aspergillus oryzae* triose phosphate isomerase, *Aspergillus nidulans* acetamidase and hybrids thereof. In a yeast host, a useful promoter is the *Saccharomyces cerevisiae* enolase (*eno-1*) promoter. Particularly preferred promoters are the TAKA amylase, NA2-tpi (a hybrid of the promoters from the genes encoding *Aspergillus niger* neutral alpha -amylase and *Aspergillus oryzae* triose phosphate isomerase) and *glaA* promoters.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be operably linked to a terminator sequence at its 3' terminus. The terminator sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any terminator which is functional in the fungal host cell of choice may be used in the present invention, but particularly preferred terminators are obtained from the genes encoding *Aspergillus oryzae* TAKA amylase, *Aspergillus niger* glucoamylase, *Aspergillus nidulans* anthranilate synthase, *Aspergillus niger* alpha-glucosidase and *Saccharomyces cerevisiae* enolase.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be operably linked to a suitable leader sequence. A leader sequence is a nontranslated region of a mRNA which is important for translation by the fungal host. The leader sequence is operably linked to the 5' terminus of the nucleic acid sequence encoding the protein or fragment thereof. The leader sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any leader sequence which is functional in the fungal host cell of choice may be used in the present invention, but particularly preferred leaders are obtained from the genes encoding *Aspergillus oryzae* TAKA amylase and *Aspergillus oryzae* triose phosphate isomerase.

A polyadenylation sequence may also be operably linked to the 3' terminus of the nucleic acid sequence of the present invention. The polyadenylation sequence is a sequence which when transcribed is recognized by the fungal host to add polyadenosine residues to transcribed mRNA. The polyadenylation sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any polyadenylation sequence which is functional in the fungal host of choice may be used in the present invention, but particularly

preferred polyadenylation sequences are obtained from the genes encoding *Aspergillus oryzae* TKA amylase, *Aspergillus niger* glucoamylase, *Aspergillus nidulans* anthranilate synthase and *Aspergillus niger* alpha-glucosidase.

To avoid the necessity of disrupting the cell to obtain the protein or fragment thereof and to minimize the amount of possible degradation of the expressed protein or fragment thereof within the cell, it is preferred that expression of the protein or fragment thereof gives rise to a product secreted outside the cell. To this end, a protein or fragment thereof of the present invention may be linked to a signal peptide linked to the amino terminus of the protein or fragment thereof. A signal peptide is an amino acid sequence which permits the secretion of the protein or fragment thereof from the fungal host into the culture medium. The signal peptide may be native to the protein or fragment thereof of the invention or may be obtained from foreign sources. The 5' end of the coding sequence of the nucleic acid sequence of the present invention may inherently contain a signal peptide coding region naturally linked in translation reading frame with the segment of the coding region which encodes the secreted protein or fragment thereof. Alternatively, the 5' end of the coding sequence may contain a signal peptide coding region which is foreign to that portion of the coding sequence which encodes the secreted protein or fragment thereof. The foreign signal peptide may be required where the coding sequence does not normally contain a signal peptide coding region. Alternatively, the foreign signal peptide may simply replace the natural signal peptide to obtain enhanced secretion of the desired protein or fragment thereof. The foreign signal peptide coding region may be obtained from a glucoamylase or an amylase gene from an *Aspergillus* species, a lipase or proteinase gene from *Rhizomucor miehei*, the gene for the alpha-factor from *Saccharomyces cerevisiae*, or the calf preprochymosin gene. An effective signal peptide for fungal host cells is the *Aspergillus*

oryzae TAKA amylase signal, *Aspergillus niger* neutral amylase signal, the *Rhizomucor miehei* aspartic proteinase signal, the *Humicola lanuginosus* cellulase signal, or the *Rhizomucor miehei* lipase signal. However, any signal peptide capable of permitting secretion of the protein or fragment thereof in a fungal host of choice may be used in the present invention.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be linked to a propeptide coding region. A propeptide is an amino acid sequence found at the amino terminus of a protein or proenzyme. Cleavage of the propeptide from the proprotein yields a mature biochemically active protein. The resulting polypeptide is known as a propeptide or proenzyme (or a zymogen in some cases). Propeptides are generally inactive and can be converted to mature active polypeptides by catalytic or autocatalytic cleavage of the propeptide from the propeptide or proenzyme. The propeptide coding region may be native to the protein or fragment thereof or may be obtained from foreign sources. The foreign propeptide coding region may be obtained from the *Saccharomyces cerevisiae* alpha-factor gene or *Myceliophthora thermophila* laccase gene (WO 95/33836, the entirety of which is herein incorporated by reference).

The procedures used to ligate the elements described above to construct the recombinant expression vector of the present invention are well known to one skilled in the art (see, for example, Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual*, 2nd ed., Cold Spring Harbor, N.Y., (1989)).

The present invention also relates to recombinant fungal host cells produced by the methods of the present invention which are advantageously used with the recombinant vector of the present invention. The cell is preferably transformed with a vector comprising a nucleic acid sequence of the invention followed by integration of the vector into the host chromosome. The

choice of fungal host cells will to a large extent depend upon the gene encoding the protein or fragment thereof and its source. The fungal host cell may, for example, be a yeast cell or a filamentous fungal cell.

"Yeast" as used herein includes *Ascosporogenous* yeast (*Endomycetales*), *Basidiosporogenous* yeast and yeast belonging to the *Fungi Imperfecti* (*Blastomycetes*). The *Ascosporogenous* yeasts are divided into the families *Spermophthoraceae* and *Saccharomycetaceae*. The latter is comprised of four subfamilies, *Schizosaccharomycoideae* (for example, genus *Schizosaccharomyces*), *Nadsonioideae*, *Lipomycoideae* and *Saccharomycoideae* (for example, genera *Pichia*, *Kluyveromyces* and *Saccharomyces*). The *Basidiosporogenous* yeasts include the genera *Leucosporidim*, *Rhodosporeidium*, *Sporidiobolus*, *Filobasidium* and *Filobasidiella*. Yeast belonging to the *Fungi Imperfecti* are divided into two families, *Sporobolomycetaceae* (for example, genera *Sorobolomyces* and *Bullera*) and *Cryptococcaceae* (for example, genus *Candida*). Since the classification of yeast may change in the future, for the purposes of this invention, yeast shall be defined as described in Biology and Activities of Yeast (Skinner *et al.*, *Soc. App. Bacteriol. Symposium Series* No. 9, (1980), the entirety of which is herein incorporated by reference). The biology of yeast and manipulation of yeast genetics are well known in the art (*see*, for example, *Biochemistry and Genetics of Yeast*, Bacil *et al.* (ed.), 2nd edition, 1987; *The Yeasts*, Rose and Harrison (eds.), 2nd ed., (1987); and *The Molecular Biology of the Yeast Saccharomyces*, Strathern *et al.* (eds.), (1981), all of which are herein incorporated by reference in their entirety).

"Fungi" as used herein includes the phyla *Ascomycota*, *Basidiomycota*, *Chytridiomycota* and *Zygomycota* (as defined by Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK; the entirety of

which is herein incorporated by reference) as well as the *Oomycota* (as cited in Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK) and all mitosporic fungi (Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK). Representative groups of *Ascomycota* include, for example, *Neurospora*, *Eupenicillium* (= *Penicillium*), *Emericella* (= *Aspergillus*), *Eurotium* (= *Aspergillus*) and the true yeasts listed above. Examples of *Basidiomycota* include mushrooms, rusts and smuts.

Representative groups of *Chytridiomycota* include, for example, *Allomyces*, *Blastocladiella*, *Coelomomyces* and aquatic fungi. Representative groups of *Oomycota* include, for example, *Saprolegniomycetous* aquatic fungi (water molds) such as *Achlya*. Examples of mitosporic fungi include *Aspergillus*, *Penicillium*, *Candida* and *Alternaria*. Representative groups of *Zygomycota* include, for example, *Rhizopus* and *Mucor*.

"Filamentous fungi" include all filamentous forms of the subdivision *Eumycota* and *Oomycota* (as defined by Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK). The filamentous fungi are characterized by a vegetative mycelium composed of chitin, cellulose, glucan, chitosan, mannan and other complex polysaccharides. Vegetative growth is by hyphal elongation and carbon catabolism is obligately aerobic. In contrast, vegetative growth by yeasts such as *Saccharomyces cerevisiae* is by budding of a unicellular thallus and carbon catabolism may be fermentative.

In one embodiment, the fungal host cell is a yeast cell. In a preferred embodiment, the yeast host cell is a cell of the species of *Candida*, *Kluyveromyces*, *Saccharomyces*, *Schizosaccharomyces*, *Pichia* and *Yarrowia*. In a preferred embodiment, the yeast host cell is a

Saccharomyces cerevisiae cell, a *Saccharomyces carlsbergensis*, *Saccharomyces diastaticus* cell, a *Saccharomyces douglasii* cell, a *Saccharomyces kluyveri* cell, a *Saccharomyces norbensis* cell, or a *Saccharomyces oviformis* cell. In another preferred embodiment, the yeast host cell is a *Kluyveromyces lactis* cell. In another preferred embodiment, the yeast host cell is a *Yarrowia lipolytica* cell.

In another embodiment, the fungal host cell is a filamentous fungal cell. In a preferred embodiment, the filamentous fungal host cell is a cell of the species of, but not limited to, *Acremonium*, *Aspergillus*, *Fusarium*, *Humicola*, *Myceliophthora*, *Mucor*, *Neurospora*, *Penicillium*, *Thielavia*, *Tolypocladium* and *Trichoderma*. In a preferred embodiment, the filamentous fungal host cell is an *Aspergillus* cell. In another preferred embodiment, the filamentous fungal host cell is an *Acremonium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Fusarium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Humicola* cell. In another preferred embodiment, the filamentous fungal host cell is a *Myceliophthora* cell. In another even preferred embodiment, the filamentous fungal host cell is a *Mucor* cell. In another preferred embodiment, the filamentous fungal host cell is a *Neurospora* cell. In another preferred embodiment, the filamentous fungal host cell is a *Penicillium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Thielavia* cell. In another preferred embodiment, the filamentous fungal host cell is a *Tolypocladium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Trichoderma* cell. In a preferred embodiment, the filamentous fungal host cell is an *Aspergillus oryzae* cell, an *Aspergillus niger* cell, an *Aspergillus foetidus* cell, or an *Aspergillus japonicus* cell. In another preferred embodiment, the filamentous fungal host cell is a *Fusarium oxysporum* cell or a *Fusarium graminearum* cell. In another preferred embodiment, the filamentous fungal

host cell is a *Humicola insolens* cell or a *Humicola lanuginosus* cell. In another preferred embodiment, the filamentous fungal host cell is a *Myceliophthora thermophila* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Mucor miehei* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Neurospora crassa* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Penicillium purpurogenum* cell. In another most preferred embodiment, the filamentous fungal host cell is a *Thielavia terrestris* cell. In another most preferred embodiment, the *Trichoderma* cell is a *Trichoderma reesei* cell, a *Trichoderma viride* cell, a *Trichoderma longibrachiatum* cell, a *Trichoderma harzianum* cell, or a *Trichoderma koningii* cell. In a preferred embodiment, the fungal host cell is selected from an *A. nidulans* cell, an *A. niger* cell, an *A. oryzae* cell and an *A. sojae* cell. In a further preferred embodiment, the fungal host cell is an *A. nidulans* cell.

The recombinant fungal host cells of the present invention may further comprise one or more sequences which encode one or more factors that are advantageous in the expression of the protein or fragment thereof, for example, an activator (e.g., a trans-acting factor), a chaperone and a processing protease. The nucleic acids encoding one or more of these factors are preferably not operably linked to the nucleic acid encoding the protein or fragment thereof. An activator is a protein which activates transcription of a nucleic acid sequence encoding a polypeptide (Kudla *et al.*, *EMBO* 9:1355-1364(1990); Jarai and Buxton, *Current Genetics* 26:2238-244(1994); Verdier, *Yeast* 6:271-297(1990), all of which are herein incorporated by reference in their entirety). The nucleic acid sequence encoding an activator may be obtained from the genes encoding *Saccharomyces cerevisiae* heme activator protein 1 (hap1), *Saccharomyces cerevisiae* galactose metabolizing protein 4 (gal4) and *Aspergillus nidulans* ammonia regulation protein (areA). For further examples, see Verdier, *Yeast* 6:271-297 (1990);

MacKenzie *et al.*, *Journal of Gen. Microbiol.* 139:2295-2307 (1993), both of which are herein incorporated by reference in their entirety). A chaperone is a protein which assists another protein in folding properly (Hartl *et al.*, *TIBS* 19:20-25 (1994); Bergeron *et al.*, *TIBS* 19:124-128 (1994); Demolder *et al.*, *J. Biotechnology* 32:179-189 (1994); Craig, *Science* 260:1902-1903(1993); Gething and Sambrook, *Nature* 355:33-45 (1992); Puig and Gilbert, *J Biol. Chem.* 269:7764-7771 (1994); Wang and Tsou, *FASEB Journal* 7:1515-11157 (1993); Robinson *et al.*, *Bio/Technology* 1:381-384 (1994), all of which are herein incorporated by reference in their entirety). The nucleic acid sequence encoding a chaperone may be obtained from the genes encoding *Aspergillus oryzae* protein disulphide isomerase, *Saccharomyces cerevisiae* calnexin, *Saccharomyces cerevisiae* BiP/GRP78 and *Saccharomyces cerevisiae* Hsp70. For further examples, see Gething and Sambrook, *Nature* 355:33-45 (1992); Hartl *et al.*, *TIBS* 19:20-25 (1994). A processing protease is a protease that cleaves a propeptide to generate a mature biochemically active polypeptide (Enderlin and Ogrydziak, *Yeast* 10:67-79 (1994); Fuller *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:1434-1438 (1989); Julius *et al.*, *Cell* 37:1075-1089 (1984); Julius *et al.*, *Cell* 32:839-852 (1983), all of which are incorporated by reference in their entirety). The nucleic acid sequence encoding a processing protease may be obtained from the genes encoding *Aspergillus niger* Kex2, *Saccharomyces cerevisiae* dipeptidylaminopeptidase, *Saccharomyces cerevisiae* Kex2 and *Yarrowia lipolytica* dibasic processing endoprotease (xpr6). Any factor that is functional in the fungal host cell of choice may be used in the present invention.

Fungal cells may be transformed by a process involving protoplast formation, transformation of the protoplasts and regeneration of the cell wall in a manner known per se. Suitable procedures for transformation of *Aspergillus* host cells are described in EP 238 023 and

Yelton *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 81:1470-1474 (1984), both of which are herein incorporated by reference in their entirety. A suitable method of transforming *Fusarium* species is described by Malardier *et al.*, *Gene* 78:147-156 (1989), the entirety of which is herein incorporated by reference. Yeast may be transformed using the procedures described by Becker and Guarente, In: Abelson and Simon, (eds.), *Guide to Yeast Genetics and Molecular Biology, Methods Enzymol.* Volume 194, pp 182-187, Academic Press, Inc., New York; Ito *et al.*, *J. Bacteriology* 153:163 (1983); Hinnen *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 75:1920 (1978), all of which are herein incorporated by reference in their entirety.

The present invention also relates to methods of producing the protein or fragment thereof comprising culturing the recombinant fungal host cells under conditions conducive for expression of the protein or fragment thereof. The fungal cells of the present invention are cultivated in a nutrient medium suitable for production of the protein or fragment thereof using methods known in the art. For example, the cell may be cultivated by shake flask cultivation, small-scale or large-scale fermentation (including continuous, batch, fed-batch, or solid state fermentations) in laboratory or industrial fermentors performed in a suitable medium and under conditions allowing the protein or fragment thereof to be expressed and/or isolated. The cultivation takes place in a suitable nutrient medium comprising carbon and nitrogen sources and inorganic salts, using procedures known in the art (*see, e.g.*, Bennett and LaSure (eds.), *More Gene Manipulations in Fungi*, Academic Press, CA, (1991), the entirety of which is herein incorporated by reference). Suitable media are available from commercial suppliers or may be prepared according to published compositions (*e.g.*, in catalogues of the American Type Culture Collection, Manassas, VA). If the protein or fragment thereof is secreted into the nutrient

medium, a protein or fragment thereof can be recovered directly from the medium. If the protein or fragment thereof is not secreted, it is recovered from cell lysates.

The expressed protein or fragment thereof may be detected using methods known in the art that are specific for the particular protein or fragment. These detection methods may include the use of specific antibodies, formation of an enzyme product, or disappearance of an enzyme substrate. For example, if the protein or fragment thereof has enzymatic activity, an enzyme assay may be used. Alternatively, if polyclonal or monoclonal antibodies specific to the protein or fragment thereof are available, immunoassays may be employed using the antibodies to the protein or fragment thereof. The techniques of enzyme assay and immunoassay are well known to those skilled in the art.

The resulting protein or fragment thereof may be recovered by methods known in the arts. For example, the protein or fragment thereof may be recovered from the nutrient medium by conventional procedures including, but not limited to, centrifugation, filtration, extraction, spray-drying, evaporation, or precipitation. The recovered protein or fragment thereof may then be further purified by a variety of chromatographic procedures, e.g., ion exchange chromatography, gel filtration chromatography, affinity chromatography, or the like.

(c) Mammalian Constructs and Transformed Mammalian Cells

The present invention also relates to methods for obtaining a recombinant mammalian host cell, comprising introducing into a mammalian host cell exogenous genetic material. The present invention also relates to a mammalian cell comprising a mammalian recombinant vector. The present invention also relates to methods for obtaining a recombinant mammalian host cell, comprising introducing into a mammalian cell exogenous genetic material.

Mammalian cell lines available as hosts for expression are known in the art and include many immortalized cell lines available from the American Type Culture Collection (ATCC, Manassas, VA), such as HeLa cells, Chinese hamster ovary (CHO) cells, baby hamster kidney (BHK) cells and a number of other cell lines. Suitable promoters for mammalian cells are also known in the art and include viral promoters such as that from Simian Virus 40 (SV40) (Fiers *et al.*, *Nature* 273:113 (1978), the entirety of which is herein incorporated by reference), Rous sarcoma virus (RSV), adenovirus (ADV) and bovine papilloma virus (BPV). Mammalian cells may also require terminator sequences and poly-A addition sequences. Enhancer sequences which increase expression may also be included and sequences which promote amplification of the gene may also be desirable (for example methotrexate resistance genes).

Vectors suitable for replication in mammalian cells may include viral replicons, or sequences which insure integration of the appropriate sequences encoding HCV epitopes into the host genome. For example, another vector used to express foreign DNA is vaccinia virus. In this case, for example, a nucleic acid molecule encoding a protein or fragment thereof is inserted into the vaccinia genome. Techniques for the insertion of foreign DNA into the vaccinia virus genome are known in the art and may utilize, for example, homologous recombination. Such heterologous DNA is generally inserted into a gene which is non-essential to the virus, for example, the thymidine kinase gene (tk), which also provides a selectable marker. Plasmid vectors that greatly facilitate the construction of recombinant viruses have been described (*see*, for example, Mackett *et al.*, *J Virol.* 49:857 (1984); Chakrabarti *et al.*, *Mol. Cell. Biol.* 5:3403 (1985); Moss, In: *Gene Transfer Vectors For Mammalian Cells* (Miller and Calos, eds., Cold Spring Harbor Laboratory, N.Y., p. 10, (1987); all of which are herein incorporated by reference

in their entirety). Expression of the HCV polypeptide then occurs in cells or animals which are infected with the live recombinant vaccinia virus.

The sequence to be integrated into the mammalian sequence may be introduced into the primary host by any convenient means, which includes calcium precipitated DNA, spheroplast fusion, transformation, electroporation, biolistics, lipofection, microinjection, or other convenient means. Where an amplifiable gene is being employed, the amplifiable gene may serve as the selection marker for selecting hosts into which the amplifiable gene has been introduced. Alternatively, one may include with the amplifiable gene another marker, such as a drug resistance marker, e.g. neomycin resistance (G418 in mammalian cells), hygromycin in resistance etc., or an auxotrophy marker (HIS3, TRP1, LEU2, URA3, ADE2, LYS2, etc.) for use in yeast cells.

Depending upon the nature of the modification and associated targeting construct, various techniques may be employed for identifying targeted integration. Conveniently, the DNA may be digested with one or more restriction enzymes and the fragments probed with an appropriate DNA fragment which will identify the properly sized restriction fragment associated with integration.

One may use different promoter sequences, enhancer sequences, or other sequence which will allow for enhanced levels of expression in the expression host. Thus, one may combine an enhancer from one source, a promoter region from another source, a 5'- noncoding region upstream from the initiation methionine from the same or different source as the other sequences and the like. One may provide for an intron in the non-coding region with appropriate splice sites or for an alternative 3'- untranslated sequence or polyadenylation site. Depending upon the particular purpose of the modification, any of these sequences may be introduced, as desired.

Where selection is intended, the sequence to be integrated will have with it a marker gene, which allows for selection. The marker gene may conveniently be downstream from the target gene and may include resistance to a cytotoxic agent, e.g. antibiotics, heavy metals, or the like, resistance or susceptibility to HAT, gancyclovir, etc., complementation to an auxotrophic host, particularly by using an auxotrophic yeast as the host for the subject manipulations, or the like. The marker gene may also be on a separate DNA molecule, particularly with primary mammalian cells. Alternatively, one may screen the various transformants, due to the high efficiency of recombination in yeast, by using hybridization analysis, PCR, sequencing, or the like.

For homologous recombination, constructs can be prepared where the amplifiable gene will be flanked, normally on both sides with DNA homologous with the DNA of the target region. Depending upon the nature of the integrating DNA and the purpose of the integration, the homologous DNA will generally be within 100kb, usually 50kb, preferably about 25kb, of the transcribed region of the target gene, more preferably within 2kb of the target gene. Where modeling of the gene is intended, homology will usually be present proximal to the site of the mutation. The homologous DNA may include the 5'-upstream region outside of the transcriptional regulatory region or comprising any enhancer sequences, transcriptional initiation sequences, adjacent sequences, or the like. The homologous region may include a portion of the coding region, where the coding region may be comprised only of an open reading frame or combination of exons and introns. The homologous region may comprise all or a portion of an intron, where all or a portion of one or more exons may also be present. Alternatively, the homologous region may comprise the 3'-region, so as to comprise all or a portion of the transcriptional termination region, or the region 3' of this region. The homologous regions may

extend over all or a portion of the target gene or be outside the target gene comprising all or a portion of the transcriptional regulatory regions and/or the structural gene.

The integrating constructs may be prepared in accordance with conventional ways, where sequences may be synthesized, isolated from natural sources, manipulated, cloned, ligated, subjected to in vitro mutagenesis, primer repair, or the like. At various stages, the joined sequences may be cloned and analyzed by restriction analysis, sequencing, or the like. Usually during the preparation of a construct where various fragments are joined, the fragments, intermediate constructs and constructs will be carried on a cloning vector comprising a replication system functional in a prokaryotic host, e.g., *E. coli* and a marker for selection, e.g., biocide resistance, complementation to an auxotrophic host, etc. Other functional sequences may also be present, such as polylinkers, for ease of introduction and excision of the construct or portions thereof, or the like. A large number of cloning vectors are available such as pBR322, the pUC series, etc. These constructs may then be used for integration into the primary mammalian host.

In the case of the primary mammalian host, a replicating vector may be used. Usually, such vector will have a viral replication system, such as SV40, bovine papilloma virus, adenovirus, or the like. The linear DNA sequence vector may also have a selectable marker for identifying transfected cells. Selectable markers include the neo gene, allowing for selection with G418, the herpes tk gene for selection with HAT medium, the gpt gene with mycophenolic acid, complementation of an auxotrophic host, etc.

The vector may or may not be capable of stable maintenance in the host. Where the vector is capable of stable maintenance, the cells will be screened for homologous integration of the vector into the genome of the host, where various techniques for curing the cells may be

employed. Where the vector is not capable of stable maintenance, for example, where a temperature sensitive replication system is employed, one may change the temperature from the permissive temperature to the non-permissive temperature, so that the cells may be cured of the vector. In this case, only those cells having integration of the construct comprising the amplifiable gene and, when present, the selectable marker, will be able to survive selection.

Where a selectable marker is present, one may select for the presence of the targeting construct by means of the selectable marker. Where the selectable marker is not present, one may select for the presence of the construct by the amplifiable gene. For the neo gene or the herpes tk gene, one could employ a medium for growth of the transformants of about 0.1-1 mg/ml of G418 or may use HAT medium, respectively. Where DHFR is the amplifiable gene, the selective medium may include from about 0.01-0.5 μ M of methotrexate or be deficient in glycine-hypoxanthine-thymidine and have dialysed serum (GHT media).

The DNA can be introduced into the expression host by a variety of techniques that include calcium phosphate/DNA co-precipitates, microinjection of DNA into the nucleus, electroporation, yeast protoplast fusion with intact cells, transfection, polycations, e.g., polybrene, polyornithine, etc., or the like. The DNA may be single or double stranded DNA, linear or circular. The various techniques for transforming mammalian cells are well known (see Keown *et al.*, *Methods Enzymol.* (1989); Keown *et al.*, *Methods Enzymol.* 185:527-537 (1990); Mansour *et al.*, *Nature* 336:348-352, (1988); all of which are herein incorporated by reference in their entirety).

(d) Insect Constructs and Transformed Insect Cells

The present invention also relates to an insect recombinant vectors comprising exogenous genetic material. The present invention also relates to an insect cell comprising an insect

recombinant vector. The present invention also relates to methods for obtaining a recombinant insect host cell, comprising introducing into an insect cell exogenous genetic material.

The insect recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures and can bring about the expression of the nucleic acid sequence. The choice of a vector will typically depend on the compatibility of the vector with the insect host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the insect host. In addition, the insect vector may be an expression vector. Nucleic acid molecules can be suitably inserted into a replication vector for expression in the insect cell under a suitable promoter for insect cells. Many vectors are available for this purpose and selection of the appropriate vector will depend mainly on the size of the nucleic acid molecule to be inserted into the vector and the particular host cell to be transformed with the vector. Each vector contains various components depending on its function (amplification of DNA or expression of DNA) and the particular host cell with which it is compatible. The vector components for insect cell transformation generally include, but are not limited to, one or more of the following: a signal sequence, origin of replication, one or more marker genes and an inducible promoter.

The insect vector may be an autonomously replicating vector, *i.e.*, a vector which exists as an extrachromosomal entity, the replication of which is independent of chromosomal replication, *e.g.*, a plasmid, an extrachromosomal element, a minichromosome, or an artificial chromosome. The vector may contain any means for assuring self-replication. Alternatively, the vector may be one which, when introduced into the insect cell, is integrated into the genome and replicated together with the chromosome(s) into which it has been integrated. For integration,

the vector may rely on the nucleic acid sequence of the vector for stable integration of the vector into the genome by homologous or nonhomologous recombination. Alternatively, the vector may contain additional nucleic acid sequences for directing integration by homologous recombination into the genome of the insect host. The additional nucleic acid sequences enable the vector to be integrated into the host cell genome at a precise location(s) in the chromosome(s). To increase the likelihood of integration at a precise location, there should be preferably two nucleic acid sequences which individually contain a sufficient number of nucleic acids, preferably 400bp to 1500bp, more preferably 800bp to 1000bp, which are highly homologous with the corresponding target sequence to enhance the probability of homologous recombination. These nucleic acid sequences may be any sequence that is homologous with a target sequence in the genome of the insect host cell and, furthermore, may be non-encoding or encoding sequences.

Baculovirus expression vectors (BEVs) have become important tools for the expression of foreign genes, both for basic research and for the production of proteins with direct clinical applications in human and veterinary medicine (Doerfler, *Curr. Top. Microbiol. Immunol.* 131:51-68 (1968); Luckow and Summers, *Bio/Technology* 6:47-55 (1988a); Miller, *Annual Review of Microbiol.* 42:177-199 (1988); Summers, *Curr. Comm. Molecular Biology*, Cold Spring Harbor Press, Cold Spring Harbor, N.Y. (1988); all of which are herein incorporated by reference in their entirety). BEVs are recombinant insect viruses in which the coding sequence for a chosen foreign gene has been inserted behind a baculovirus promoter in place of the viral gene, e.g., polyhedrin (Smith and Summers, U.S. Pat. No., 4,745,051, the entirety of which is incorporated herein by reference).

The use of baculovirus vectors relies upon the host cells being derived from *Lepidopteran* insects such as *Spodoptera frugiperda* or *Trichoplusia ni*. The preferred *Spodoptera frugiperda* cell line is the cell line Sf9. The *Spodoptera frugiperda* Sf9 cell line was obtained from American Type Culture Collection (Manassas, VA.) and is assigned accession number ATCC CRL 1711 (Summers and Smith, *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Ag. Exper. Station Bulletin No. 1555 (1988), the entirety of which is herein incorporated by reference). Other insect cell systems, such as the silkworm *B. mori* may also be used.

The proteins expressed by the BEVs are, therefore, synthesized, modified and transported in host cells derived from *Lepidopteran* insects. Most of the genes that have been inserted and produced in the baculovirus expression vector system have been derived from vertebrate species. Other baculovirus genes in addition to the polyhedrin promoter may be employed to advantage in a baculovirus expression system. These include immediate-early (alpha), delayed-early (β), late (γ), or very late (delta), according to the phase of the viral infection during which they are expressed. The expression of these genes occurs sequentially, probably as the result of a "cascade" mechanism of transcriptional regulation. (Guarino and Summers, *J. Virol.* 57:563-571 (1986); Guarino and Summers, *J. Virol.* 61:2091-2099 (1987); Guarino and Summers, *Virol.* 162:444-451 (1988); all of which are herein incorporated by reference in their entirety).

Insect recombinant vectors are useful as intermediates for the infection or transformation of insect cell systems. For example, an insect recombinant vector containing a nucleic acid molecule encoding a baculovirus transcriptional promoter followed downstream by an insect signal DNA sequence is capable of directing the secretion of the desired biologically active protein from the insect cell. The vector may utilize a baculovirus transcriptional promoter region

derived from any of the over 500 baculoviruses generally infecting insects, such as for example the Orders *Lepidoptera*, *Diptera*, *Orthoptera*, *Coleoptera* and *Hymenoptera*, including for example but not limited to the viral DNAs of *Autographa californica* MNPV, *Bombyx mori* NPV, *Trichoplusia ni* MNPV, *Rachiplusia ou* MNPV or *Galleria mellonella* MNPV, wherein said baculovirus transcriptional promoter is a baculovirus immediate-early gene IEl or IEN promoter; an immediate-early gene in combination with a baculovirus delayed-early gene promoter region selected from the group consisting of 39K and a *HindIII-k* fragment delayed-early gene; or a baculovirus late gene promoter. The immediate-early or delayed-early promoters can be enhanced with transcriptional enhancer elements. The insect signal DNA sequence may code for a signal peptide of a *Lepidopteran* adipokinetic hormone precursor or a signal peptide of the *Manduca sexta* adipokinetic hormone precursor (Summers, U.S. Patent No. 5,155,037; the entirety of which is herein incorporated by reference). Other insect signal DNA sequences include a signal peptide of the *Orthoptera Schistocerca gregaria* locust adipokinetic hormone precursor and the *Drosophila melanogaster* cuticle genes CP1, CP2, CP3 or CP4 or for an insect signal peptide having substantially a similar chemical composition and function (Summers, U.S. Patent No. 5,155,037).

Insect cells are distinctly different from animal cells. Insects have a unique life cycle and have distinct cellular properties such as the lack of intracellular plasminogen activators in which are present in vertebrate cells. Another difference is the high expression levels of protein products ranging from 1 to greater than 500 mg/liter and the ease at which cDNA can be cloned into cells (Frasier, *In Vitro Cell. Dev. Biol.* 25:225 (1989); Summers and Smith, In: *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Ag. Exper. Station Bulletin No. 1555 (1988), both of which are incorporated by reference in their entirety).

Recombinant protein expression in insect cells is achieved by viral infection or stable transformation. For viral infection, the desired gene is cloned into baculovirus at the site of the wild-type polyhedron gene (Webb and Summers, *Technique* 2:173 (1990); Bishop and Posse, *Adv. Gene Technol.* 1:55 (1990); both of which are incorporated by reference in their entirety). The polyhedron gene is a component of a protein coat in occlusions which encapsulate virus particles. Deletion or insertion in the polyhedron gene results the failure to form occlusion bodies. Occlusion negative viruses are morphologically different from occlusion positive viruses and enable one skilled in the art to identify and purify recombinant viruses.

The vectors of present invention preferably contain one or more selectable markers which permit easy selection of transformed cells. A selectable marker is a gene the product of which provides, for example biocide or viral resistance, resistance to heavy metals, prototrophy to auxotrophs and the like. Selection may be accomplished by co-transformation, *e.g.*, as described in WO 91/17243, a nucleic acid sequence of the present invention may be operably linked to a suitable promoter sequence. The promoter sequence is a nucleic acid sequence which is recognized by the insect host cell for expression of the nucleic acid sequence. The promoter sequence contains transcription and translation control sequences which mediate the expression of the protein or fragment thereof. The promoter may be any nucleic acid sequence which shows transcriptional activity in the insect host cell of choice and may be obtained from genes encoding polypeptides either homologous or heterologous to the host cell.

For example, a nucleic acid molecule encoding a protein or fragment thereof may also be operably linked to a suitable leader sequence. A leader sequence is a nontranslated region of a mRNA which is important for translation by the fungal host. The leader sequence is operably linked to the 5' terminus of the nucleic acid sequence encoding the protein or fragment thereof.

The leader sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any leader sequence which is functional in the insect host cell of choice may be used in the present invention.

A polyadenylation sequence may also be operably linked to the 3' terminus of the nucleic acid sequence of the present invention. The polyadenylation sequence is a sequence which when transcribed is recognized by the insect host to add polyadenosine residues to transcribed mRNA. The polyadenylation sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any polyadenylation sequence which is functional in the fungal host of choice may be used in the present invention.

To avoid the necessity of disrupting the cell to obtain the protein or fragment thereof and to minimize the amount of possible degradation of the expressed polypeptide within the cell, it is preferred that expression of the polypeptide gene gives rise to a product secreted outside the cell. To this end, the protein or fragment thereof of the present invention may be linked to a signal peptide linked to the amino terminus of the protein or fragment thereof. A signal peptide is an amino acid sequence which permits the secretion of the protein or fragment thereof from the insect host into the culture medium. The signal peptide may be native to the protein or fragment thereof of the invention or may be obtained from foreign sources. The 5' end of the coding sequence of the nucleic acid sequence of the present invention may inherently contain a signal peptide coding region naturally linked in translation reading frame with the segment of the coding region which encodes the secreted protein or fragment thereof.

At present, a mode of achieving secretion of a foreign gene product in insect cells is by way of the foreign gene's native signal peptide. Because the foreign genes are usually from non-insect organisms, their signal sequences may be poorly recognized by insect cells and hence,

levels of expression may be suboptimal. However, the efficiency of expression of foreign gene products seems to depend primarily on the characteristics of the foreign protein. On average, nuclear localized or non-structural proteins are most highly expressed, secreted proteins are intermediate and integral membrane proteins are the least expressed. One factor generally affecting the efficiency of the production of foreign gene products in a heterologous host system is the presence of native signal sequences (also termed presequences, targeting signals, or leader sequences) associated with the foreign gene. The signal sequence is generally coded by a DNA sequence immediately following (5' to 3') the translation start site of the desired foreign gene.

The expression dependence on the type of signal sequence associated with a gene product can be represented by the following example: If a foreign gene is inserted at a site downstream from the translational start site of the baculovirus polyhedrin gene so as to produce a fusion protein (containing the N-terminus of the polyhedrin structural gene), the fused gene is highly expressed. But less expression is achieved when a foreign gene is inserted in a baculovirus expression vector immediately following the transcriptional start site and totally replacing the polyhedrin structural gene.

Insertions into the region -50 to -1 significantly alter (reduce) steady state transcription which, in turn, reduces translation of the foreign gene product. Use of the pVL941 vector optimizes transcription of foreign genes to the level of the polyhedrin gene transcription. Even though the transcription of a foreign gene may be optimal, optimal translation may vary because of several factors involving processing: signal peptide recognition, mRNA and ribosome binding, glycosylation, disulfide bond formation, sugar processing, oligomerization, for example.

The properties of the insect signal peptide are expected to be more optimal for the efficiency of the translation process in insect cells than those from vertebrate proteins. This

phenomenon can generally be explained by the fact that proteins secreted from cells are synthesized as precursor molecules containing hydrophobic N-terminal signal peptides. The signal peptides direct transport of the select protein to its target membrane and are then cleaved by a peptidase on the membrane, such as the endoplasmic reticulum, when the protein passes through it.

Another exemplary insect signal sequence is the sequence encoding for *Drosophila* cuticle proteins such as CP1, CP2, CP3 or CP4 (Summers, U.S. Patent No. 5,278,050; the entirety of which is herein incorporated by reference). Most of a 9kb region of the *Drosophila* genome containing genes for the cuticle proteins has been sequenced. Four of the five cuticle genes contains a signal peptide coding sequence interrupted by a short intervening sequence (about 60 base pairs) at a conserved site. Conserved sequences occur in the 5' mRNA untranslated region, in the adjacent 35 base pairs of upstream flanking sequence and at -200 base pairs from the mRNA start position in each of the cuticle genes.

Standard methods of insect cell culture, cotransfection and preparation of plasmids are set forth in Summers and Smith (Summers and Smith, *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Agricultural Experiment Station Bulletin No. 1555, Texas A&M University (1987)). Procedures for the cultivation of viruses and cells are described in Volkman and Summers, *J. Virol* 19:820-832 (1975) and Volkman *et al.*, *J. Virol* 19:820-832 (1976); both of which are herein incorporated by reference in their entirety.

(e) Bacterial Constructs and Transformed Bacterial Cells

The present invention also relates to a bacterial recombinant vector comprising exogenous genetic material. The present invention also relates to a bacteria cell comprising a bacterial recombinant vector. The present invention also relates to methods for obtaining a

recombinant bacteria host cell, comprising introducing into a bacterial host cell exogenous genetic material.

The bacterial recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures. The choice of a vector will typically depend on the compatibility of the vector with the bacterial host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the bacterial host. In addition, the bacterial vector may be an expression vector. Nucleic acid molecules encoding protein homologues or fragments thereof can, for example, be suitably inserted into a replicable vector for expression in the bacterium under the control of a suitable promoter for bacteria. Many vectors are available for this purpose and selection of the appropriate vector will depend mainly on the size of the nucleic acid to be inserted into the vector and the particular host cell to be transformed with the vector. Each vector contains various components depending on its function (amplification of DNA or expression of DNA) and the particular host cell with which it is compatible. The vector components for bacterial transformation generally include, but are not limited to, one or more of the following: a signal sequence, an origin of replication, one or more marker genes and an inducible promoter.

In general, plasmid vectors containing replicon and control sequences that are derived from species compatible with the host cell are used in connection with bacterial hosts. The vector ordinarily carries a replication site, as well as marking sequences that are capable of providing phenotypic selection in transformed cells. For example, *E. coli* is typically transformed using pBR322, a plasmid derived from an *E. coli* species (see, e.g., Bolivar *et al.*,

Gene 2:95 (1977); the entirety of which is herein incorporated by reference). pBR322 contains genes for ampicillin and tetracycline resistance and thus provides easy means for identifying transformed cells. The pBR322 plasmid, or other microbial plasmid or phage, also generally contains, or is modified to contain, promoters that can be used by the microbial organism for expression of the selectable marker genes.

Nucleic acid molecules encoding protein or fragments thereof may be expressed not only directly, but also as a fusion with another polypeptide, preferably a signal sequence or other polypeptide having a specific cleavage site at the N-terminus of the mature polypeptide. In general, the signal sequence may be a component of the vector, or it may be a part of the polypeptide DNA that is inserted into the vector. The heterologous signal sequence selected should be one that is recognized and processed (i.e., cleaved by a signal peptidase) by the host cell. For bacterial host cells that do not recognize and process the native polypeptide signal sequence, the signal sequence is substituted by a bacterial signal sequence selected, for example, from the group consisting of the alkaline phosphatase, penicillinase, lpp, or heat-stable enterotoxin II leaders.

Both expression and cloning vectors contain a nucleic acid sequence that enables the vector to replicate in one or more selected host cells. Generally, in cloning vectors this sequence is one that enables the vector to replicate independently of the host chromosomal DNA and includes origins of replication or autonomously replicating sequences. Such sequences are well known for a variety of bacteria. The origin of replication from the plasmid pBR322 is suitable for most Gram-negative bacteria.

Expression and cloning vectors also generally contain a selection gene, also termed a selectable marker. This gene encodes a protein necessary for the survival or growth of

transformed host cells grown in a selective culture medium. Host cells not transformed with the vector containing the selection gene will not survive in the culture medium. Typical selection genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g., ampicillin, neomycin, methotrexate, or tetracycline, (b) complement auxotrophic deficiencies, or (c) supply critical nutrients not available from complex media, e.g., the gene encoding D-alanine racemase for *Bacilli*. One example of a selection scheme utilizes a drug to arrest growth of a host cell. Those cells that are successfully transformed with a heterologous protein homologue or fragment thereof produce a protein conferring drug resistance and thus survive the selection regimen.

The expression vector for producing a protein or fragment thereof can also contains an inducible promoter that is recognized by the host bacterial organism and is operably linked to the nucleic acid encoding, for example, the nucleic acid molecule encoding the protein homologue or fragment thereof of interest. Inducible promoters suitable for use with bacterial hosts include the β -lactamase and lactose promoter systems (Chang *et al.*, *Nature* 275:615 (1978); Goeddel *et al.*, *Nature* 281:544 (1979); both of which are herein incorporated by reference in their entirety), the arabinose promoter system (Guzman *et al.*, *J. Bacteriol.* 174:7716-7728 (1992); the entirety of which is herein incorporated by reference), alkaline phosphatase, a tryptophan (trp) promoter system (Goeddel, *Nucleic Acids Res.* 8:4057 (1980); EP 36,776; both of which are herein incorporated by reference in their entirety) and hybrid promoters such as the tac promoter (deBoer *et al.*, *Proc. Natl. Acad. Sci. (USA)* 80:21-25 (1983); the entirety of which is herein incorporated by reference). However, other known bacterial inducible promoters are suitable (Siebenlist *et al.*, *Cell* 20:269 (1980); the entirety of which is herein incorporated by reference).

Promoters for use in bacterial systems also generally contain a Shine-Dalgarno (S.D.) sequence operably linked to the DNA encoding the polypeptide of interest. The promoter can be

removed from the bacterial source DNA by restriction enzyme digestion and inserted into the vector containing the desired DNA.

Construction of suitable vectors containing one or more of the above-listed components employs standard ligation techniques. Isolated plasmids or DNA fragments are cleaved, tailored and re-ligated in the form desired to generate the plasmids required. Examples of available bacterial expression vectors include, but are not limited to, the multifunctional *E. coli* cloning and expression vectors such as Bluescript™ (Stratagene, La Jolla, CA), in which, for example, encoding an *A. nidulans* protein homologue or fragment thereof homologue, may be ligated into the vector in frame with sequences for the amino-terminal Met and the subsequent 7 residues of β -galactosidase so that a hybrid protein is produced; pIN vectors (Van Heeke and Schuster, *J. Biol. Chem.* 264:5503-5509 (1989), the entirety of which is herein incorporated by reference); and the like. pGEX vectors (Promega, Madison Wisconsin U.S.A.) may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption to glutathione-agarose beads followed by elution in the presence of free glutathione. Proteins made in such systems are designed to include heparin, thrombin or factor XA protease cleavage sites so that the cloned polypeptide of interest can be released from the GST moiety at will.

Suitable host bacteria for a bacterial vector include archaebacteria and eubacteria, especially eubacteria and most preferably *Enterobacteriaceae*. Examples of useful bacteria include *Escherichia*, *Enterobacter*, *Azotobacter*, *Erwinia*, *Bacillus*, *Pseudomonas*, *Klebsiella*, *Proteus*, *Salmonella*, *Serratia*, *Shigella*, *Rhizobia*, *Vitreoscilla* and *Paracoccus*. Suitable *E. coli* hosts include *E. coli* W3110 (American Type Culture Collection (ATCC) 27,325, Manassas,

Virginia U.S.A.), *E. coli* 294 (ATCC 31,446), *E. coli* B and *E. coli* X1776 (ATCC 31,537).

These examples are illustrative rather than limiting. Mutant cells of any of the above-mentioned bacteria may also be employed. It is, of course, necessary to select the appropriate bacteria taking into consideration replicability of the replicon in the cells of a bacterium. For example, *E. coli*, *Serratia*, or *Salmonella* species can be suitably used as the host when well known plasmids such as pBR322, pBR325, pACYC177, or pKN410 are used to supply the replicon. *E. coli* strain W3110 is a preferred host or parent host because it is a common host strain for recombinant DNA product fermentations. Preferably, the host cell should secrete minimal amounts of proteolytic enzymes.

Host cells are transfected and preferably transformed with the above-described vectors and cultured in conventional nutrient media modified as appropriate for inducing promoters, selecting transformants, or amplifying the genes encoding the desired sequences.

Numerous methods of transfection are known to the ordinarily skilled artisan, for example, calcium phosphate and electroporation. Depending on the host cell used, transformation is done using standard techniques appropriate to such cells. The calcium treatment employing calcium chloride, as described in section 1.82 of Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, New York: Cold Spring Harbor Laboratory Press, (1989), is generally used for bacterial cells that contain substantial cell-wall barriers. Another method for transformation employs polyethylene glycol/DMSO, as described in Chung and Miller (Chung and Miller, *Nucleic Acids Res.* 16:3580 (1988); the entirety of which is herein incorporated by reference). Yet another method is the use of the technique termed electroporation.

Bacterial cells used to produce the polypeptide of interest for purposes of this invention are cultured in suitable media in which the promoters for the nucleic acid encoding the heterologous polypeptide can be artificially induced as described generally, e.g., in Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, New York: Cold Spring Harbor Laboratory Press, (1989). Examples of suitable media are given in U.S. Pat. Nos. 5,304,472 and 5,342,763; both of which are incorporated by reference in their entirety.

In addition to the above discussed procedures, practitioners are familiar with the standard resource materials which describe specific conditions and procedures for the construction, manipulation and isolation of macromolecules (e.g., DNA molecules, plasmids, etc.), generation of recombinant organisms and the screening and isolating of clones, (see for example, Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Press (1989); Mailga *et al.*, *Methods in Plant Molecular Biology*, Cold Spring Harbor Press (1995), the entirety of which is herein incorporated by reference; Birren *et al.*, *Genome Analysis: Analyzing DNA*, 1, Cold Spring Harbor, New York, the entirety of which is herein incorporated by reference).

(f) Computer Readable Media

The nucleotide sequence provided in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof, or complement thereof, or a nucleotide sequence at least 90% identical, preferably 95%, identical even more preferably 99% or 100% identical to the sequence provided in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof, or complement thereof, can be “provided” in a variety of mediums to facilitate use. Such a medium can also provide a subset thereof in a form that allows a skilled artisan to examine the sequences.

A preferred subset of nucleotide sequences are those nucleic acid sequences that encode a maize or soybean methionine adenosyltransferase enzyme or complement thereof or fragment of

either, a nucleic acid molecule that encodes a maize or soybean S-adenosylmethionine
 decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that
 encodes a maize or soybean aspartate kinase enzyme or complement thereof or fragment of
 either, a nucleic acid molecule that encodes a maize or soybean aspartate-semialdehyde
 dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that
 encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof
 or fragment of either, a nucleic acid molecule that encodes a maize or soybean cystathionine β -
 lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes
 a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase
 enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a
 maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of
 either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase
 enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a
 maize or a soybean cystathionine γ -lyase enzyme or complement thereof or fragment of either
 and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase
 enzyme or complement thereof or fragment of either.

A further preferred subset of nucleic acid sequences is where the subset of sequences is
 two proteins or fragments thereof, more preferably three proteins or fragments thereof and even
 more preferable four proteins or fragments thereof, these nucleic acid sequences are selected
 from the group that comprises a maize or soybean methionine adenosyltransferase enzyme or
 complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or
 soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of
 either, a nucleic acid molecule that encodes a maize or soybean aspartate kinase enzyme or

complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean cystathionine β -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine β -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine γ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either.

In one application of this embodiment, a nucleotide sequence of the present invention can be recorded on computer readable media. As used herein, "computer readable media" refers to any medium that can be read and accessed directly by a computer. Such media include, but are not limited to: magnetic storage media, such as floppy discs, hard disc, storage medium and magnetic tape; optical storage media such as CD-ROM; electrical storage media such as RAM and ROM; and hybrids of these categories such as magnetic/optical storage media. A skilled artisan can readily appreciate how any of the presently known computer readable mediums can be used to create a manufacture comprising computer readable medium having recorded thereon a nucleotide sequence of the present invention.

As used herein, "recorded" refers to a process for storing information on computer readable medium. A skilled artisan can readily adopt any of the presently known methods for recording information on computer readable medium to generate media comprising the nucleotide sequence information of the present invention. A variety of data storage structures are available to a skilled artisan for creating a computer readable medium having recorded thereon a nucleotide sequence of the present invention. The choice of the data storage structure will generally be based on the means chosen to access the stored information. In addition, a variety of data processor programs and formats can be used to store the nucleotide sequence information of the present invention on computer readable medium. The sequence information can be represented in a word processing text file, formatted in commercially-available software such as WordPerfect and Microsoft Word, or represented in the form of an ASCII file, stored in a database application, such as DB2, Sybase, Oracle, or the like. A skilled artisan can readily adapt any number of data processor structuring formats (e.g. text file or database) in order to obtain computer readable medium having recorded thereon the nucleotide sequence information of the present invention.

By providing one or more of nucleotide sequences of the present invention, a skilled artisan can routinely access the sequence information for a variety of purposes. Computer software is publicly available which allows a skilled artisan to access sequence information provided in a computer readable medium. The examples which follow demonstrate how software which implements the BLAST (Altschul *et al.*, *J. Mol. Biol.* 215:403-410 (1990), the entirety of which is herein incorporated by reference) and BLAZE (Brutlag *et al.*, *Comp. Chem.* 17:203-207 (1993), the entirety of which is herein incorporated by reference) search algorithms on a Sybase system can be used to identify open reading frames (ORFs) within the genome that

contain homology to ORFs or proteins from other organisms. Such ORFs are protein-encoding fragments within the sequences of the present invention and are useful in producing commercially important proteins such as enzymes used in amino acid biosynthesis, metabolism, transcription, translation, RNA processing, nucleic acid and a protein degradation, protein modification and DNA replication, restriction, modification, recombination and repair.

The present invention further provides systems, particularly computer-based systems, which contain the sequence information described herein. Such systems are designed to identify commercially important fragments of the nucleic acid molecule of the present invention. As used herein, “a computer-based system” refers to the hardware means, software means and data storage means used to analyze the nucleotide sequence information of the present invention. The minimum hardware means of the computer-based systems of the present invention comprises a central processing unit (CPU), input means, output means and data storage means. A skilled artisan can readily appreciate that any one of the currently available computer-based system are suitable for use in the present invention.

As indicated above, the computer-based systems of the present invention comprise a data storage means having stored therein a nucleotide sequence of the present invention and the necessary hardware means and software means for supporting and implementing a search means. As used herein, “data storage means” refers to memory that can store nucleotide sequence information of the present invention, or a memory access means which can access manufactures having recorded thereon the nucleotide sequence information of the present invention. As used herein, “search means” refers to one or more programs which are implemented on the computer-based system to compare a target sequence or target structural motif with the sequence information stored within the data storage means. Search means are used to identify fragments

or regions of the sequence of the present invention that match a particular target sequence or target motif. A variety of known algorithms are disclosed publicly and a variety of commercially available software for conducting search means are available can be used in the computer-based systems of the present invention. Examples of such software include, but are not limited to, MacPattern (EMBL), BLASTIN and BLASTIX (NCBIA). One of the available algorithms or implementing software packages for conducting homology searches can be adapted for use in the present computer-based systems.

The most preferred sequence length of a target sequence is from about 10 to 100 amino acids or from about 30 to 300 nucleotide residues. However, it is well recognized that during searches for commercially important fragments of the nucleic acid molecules of the present invention, such as sequence fragments involved in gene expression and protein processing, may be of shorter length.

As used herein, "a target structural motif," or "target motif," refers to any rationally selected sequence or combination of sequences in which the sequences the sequence(s) are chosen based on a three-dimensional configuration which is formed upon the folding of the target motif. There are a variety of target motifs known in the art. Protein target motifs include, but are not limited to, enzymatic active sites and signal sequences. Nucleic acid target motifs include, but are not limited to, promoter sequences, *cis* elements, hairpin structures and inducible expression elements (protein binding sequences).

Thus, the present invention further provides an input means for receiving a target sequence, a data storage means for storing the target sequences of the present invention sequence identified using a search means as described above and an output means for outputting the identified homologous sequences. A variety of structural formats for the input and output means

can be used to input and output information in the computer-based systems of the present invention. A preferred format for an output means ranks fragments of the sequence of the present invention by varying degrees of homology to the target sequence or target motif. Such presentation provides a skilled artisan with a ranking of sequences which contain various amounts of the target sequence or target motif and identifies the degree of homology contained in the identified fragment.

A variety of comparing means can be used to compare a target sequence or target motif with the data storage means to identify sequence fragments sequence of the present invention. For example, implementing software which implement the BLAST and BLAZE algorithms (Altschul *et al.*, *J. Mol. Biol.* 215:403-410 (1990)) can be used to identify open frames within the nucleic acid molecules of the present invention. A skilled artisan can readily recognize that any one of the publicly available homology search programs can be used as the search means for the computer-based systems of the present invention.

Having now generally described the invention, the same will be more readily understood through reference to the following examples which are provided by way of illustration and are not intended to be limiting of the present invention, unless specified.

Example 1

The MONN01 cDNA library is a normalized library generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting

at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON001 cDNA library is generated from maize (B73, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) immature tassels at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V6 stage. At that stage the tassel is an immature tassel of about 2-3 cm in length. The tassels are removed and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON003 library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) roots at the V6 developmental stage. Seeds are planted at a depth of approximately 3 cm in coil into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth, the seedlings are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and approximately 3 times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting at a concentration of 150 ppm N. Two to three times during the life time of the plant from transplanting to flowering a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in approximately 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6 leaf development stage. The root system is cut from maize plant and washed with water to free it from the soil. The tissue is then immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON004 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is

approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON005 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) root tissue at the V6 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen and the harvested tissue is then stored at -80°C until RNA preparation.

The SATMON006 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) total leaf tissue at the V6 plant development

stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON007 cDNA library is generated from the primary root tissue of 5 day old maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). After germination, the trays, along with the moist paper, are moved to a greenhouse where the maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles for approximately 5 days. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. The primary root tissue is collected when the seedlings are 5 days old. At this stage, the primary root (radicle) is

pushed through the coleorhiza which itself is pushed through the seed coat. The primary root, which is about 2-3 cm long, is cut and immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON008 cDNA library is generated from the primary shoot (coleoptile 2-3 cm) of maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings which are approximately 5 days old. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). Then the trays containing the seeds are moved to a greenhouse at 15hr daytime/9 hr nighttime cycles and grown until they are 5 days post germination. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Tissue is collected when the seedlings are 5 days old. At this stage, the primary shoot (coleoptile) is pushed through the seed coat and is about 2-3 cm long. The coleoptile is dissected away from the rest of the seedling, immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON009 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaves at the 8 leaf stage (V8 plant development stage). Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is

70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 8-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical, are cut at the base of the leaves. The leaves are then pooled and then immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON010 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) root tissue at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the V8 development stage. The root system is cut from this mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON011 cDNA library is generated from undeveloped maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaf at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium.

After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The second youngest leaf which is at the base of the apical leaf of V6 stage maize plant is cut at the base and immediately transferred to liquid nitrogen containers in which the leaf is crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON012 cDNA library is generated from 2 day post germination maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). Then the trays containing the seeds are moved to the greenhouse and grown at 15hr daytime/9 hr nighttime cycles until 2 days post germination. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Tissue is collected when the seedlings are 2 days old. At the two day stage, the coleorhiza is pushed through the seed coat and the primary root (the radicle) is pierced the coleorhiza but is barely visible. Also, at this two day stage, the coleoptile is just emerging from the seed coat. The 2 days post germination seedlings are then immersed in liquid nitrogen and crushed. The harvested tissue is stored at -80°C until preparation of total RNA.

The SATMON013 cDNA library is generated from apical maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) meristem founder at the V4 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, the plant is at the 4 leaf stage. The lead at the apex of the V4 stage maize plant is referred to as the meristem founder. This apical meristem founder is cut, immediately frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON014 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) endosperm fourteen days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9

hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the maize plant ear shoots are ready for fertilization. At this stage, the ear shoots are enclosed in a paper bag before silk emergence to withhold the pollen. The ear shoots are pollinated and 14 days after pollination, the ears are pulled out and then the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the endosperms are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON016 library is a maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) sheath library collected at the V8 developmental stage. Seeds are planted in a depth of approximately 3 cm in solid into 2-3 inch pots containing Metro growing medium. After 2-3 weeks growth, they are transplanted into 10" pots containing the same. Plants are watered daily before transplantation and approximately the times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting, at a strength of 150 ppm N. Two to three times during the life time of the plant from transplanting to flowering, a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. When the maize plants are at the V8 stage the 5th and 6th leaves from the bottom exhibit fully developed leaf blades. At the base of these leaves, the ligule is differentiated and the leaf blade is joined to the sheath. The sheath is dissected away from the

base of the leaf then the sheath is frozen in liquid nitrogen and crushed. The tissue is then stored at -80°C until RNA preparation.

The SATMON017 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) embryo seventeen days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth the seeds are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergence to withhold the pollen. The ear shoots are fertilized and 21 days after pollination, the ears are pulled out and the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the embryos are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON019 (Lib3054) cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) culm (stem) at the V8 developmental stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing

medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. When the maize plant is at the V8 stage, the 5th and 6th leaves from the bottom have fully developed leaf blades. The region between the nodes of the 5th and the sixth leaves from the bottom is the region of the stem that is collected. The leaves are pulled out and the sheath is also torn away from the stem. This stem tissue is completely free of any leaf and sheath tissue. The stem tissue is then frozen in liquid nitrogen and stored at -80°C until RNA preparation.

The SATMON020 cDNA library is from a maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) Hill Type II-Initiated Callus. Petri plates containing approximately 25 ml of Type II initiation media are prepared. This medium contains N6 salts and vitamins, 3% sucrose, 2.3 g/liter proline 0.1 g/liter enzymatic casein hydrolysate, 2mg/liter 2,4 – dichloro phenoxy-acetic acid (2,4, D), 15.3 mg/liter AgNO₃ and 0.8% bacto agar and is adjusted to pH 6.0 before autoclaving. At 9-11 days after pollination, an ear with immature embryos measuring approximately 1-2 mm in length is chosen. The husks and silks are removed and then the ear is broken into halves and placed in an autoclaved solution of Clorox/TWEEN 20 sterilizing solution. Then the ear is rinsed with deionized water. Then each embryo is extracted from the kernel. Intact embryos are placed in contact with the medium, scutellar side up). Multiple embryos are plated on each plate and the plates are incubated in the dark at 25°C. Type II

calluses are friable, can be subcultured with a spatula, frequently regenerate via somatic embryogenesis and are relatively undifferentiated. As seen in the microscope, the Tape II calluses show color ranging from translucent to light yellow and heterogeneity on with respect to embryoid structure as well as stage of embryoid development. Once Type II callus are formed, the calluses is transferred to type II callus maintenance medium without AgNO_3 . Every 7-10 days, the callus is subcultured. About 4 weeks after embryo isolation the callus is removed from the plates and then frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The SATMON021 cDNA library is generated from the immature maize (DK604, Dekalb Genetics, Dekalb Illinois, U.S.A.) tassel at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F . Supplemental lighting is provided by 1000 W sodium vapor lamps. As the maize plant enters the V8 stage, tassels which are 15-20 cm in length are collected and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The SATMON022 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) ear (growing silks) at the V8 plant development stage. Seeds are planted

at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. *Zea mays* plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the plant is in the V8 stage. At this stage, some immature ear shoots are visible. The immature ear shoots (approximately 1 cm in length) are pulled out, frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON23 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) ear (growing silk) at the V8 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. When the tissue is harvested at the V8 stage, the length of the ear that is harvested is about 10-15 cm and the silks are just exposed (approximately 1 inch).

The ear along with the silks is frozen in liquid nitrogen and then the tissue is stored at -80°C until RNA preparation.

The SATMON024 cDNA library is generated from the immature maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) tassel at the V9 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. As a maize plant enters the V9 stage, the tassel is rapidly developing and a 37 cm tassel along with the glume, anthers and pollen is collected and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The SATMON025 cDNA library is from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) Hill Type II-Regenerated Callus. Type II callus is grown in initiation media as described for SATMON020 and then the embryoids on the surface of the Type II callus are allowed to mature and germinate. The 1-2 gm fresh weight of the soft friable type callus containing numerous embryoids are transferred to 100 x 15 mm petri plates containing 25 ml of regeneration media. Regeneration media consists of Murashige and Skoog (MS) basal salts, modified White's vitamins (0.2 g/liter glycine and 0.5 g/liter myo-inositol and 0.8% bacto agar (6SMS0D)). The plates are then placed in the dark after covering with parafilm. After 1 week,

the plates are moved to a lighted growth chamber with 16 hr light and 8 hr dark photoperiod. Three weeks after plating the Type II callus to 6SMS0D, the callus exhibit shoot formation. The callus and the shoots are transferred to fresh 6SMS0D plates for another 2 weeks. The callus and the shoots are then transferred to petri plates with reduced sucrose (3SMS0D). Upon distinct formation of a root and shoot, the newly developed green plants are then removed out with a spatula and frozen in liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON026 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) juvenile/adult shift leaves at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F . Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plants are at the 8-leaf development stage. Leaves are founded sequentially around the meristem over weeks of time and the older, more juvenile leaves arise earlier and in a more basal position than the younger, more adult leaves, which are in a more apical position. In a V8 plant, some leaves which are in the middle portion of the plant exhibit characteristics of both juvenile as well as adult leaves. They exhibit a yellowing color

but also exhibit, in part, a green color. These leaves are termed juvenile/adult shift leaves. The juvenile/adult shift leaves (the 4th, 5th leaves from the bottom) are cut at the base, pooled and transferred to liquid nitrogen in which they are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON027 cDNA library is generated from 6 day maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaves. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. *Zea mays* plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, when the plant is at the 8-leaf stage, water is held back for six days. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical, are all cut at the base of the leaves. All the leaves exhibit significant wilting. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON028 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) roots at the V8 developmental stage that are subject to six days water

stress. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, when the plant is at the 8-leaf stage, water is held back for six days. The root system is cut, shaken and washed to remove soil. Root tissue is then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON029 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings at the etiolated stage. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark for 4 days at approximately 70°F. Tissue is collected when the seedlings are 4 days old. By 4 days, the primary root has penetrated the coleorhiza and is about 4-5 cm and the secondary lateral roots have also made their appearance. The coleoptile has also pushed through the seed coat and is about 4-5 cm long. The seedlings are frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON030 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) root tissue at the V4 plant development stage. Seeds are planted at a

depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth, they are transplanted into 10 inch pots containing the same. Plants are watered daily before transplantation and approximately 3 times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting, at a strength of 150 ppm N. Two to three times during the life time of the plant, from transplanting to flowering, a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 sodium vapor lamps. Tissue is collected when the maize plant is at the 4 leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is then immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON031 cDNA library is generated from the maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaf tissue at the V4 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected

when the maize plant is at the 4-leaf development stage. The third leaf from the bottom is cut at the base and immediately frozen in liquid nitrogen and crushed. The tissue is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON033 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) embryo tissue 13 days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of the maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergent to withhold the pollen. The ear shoots are pollinated and 13 days after pollination, the ears are pulled out and then the kernels are plucked cut of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the embryos are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON034 cDNA library is generated from cold stressed maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept on at 10°C for 7 days. After 7 days, the temperature is shifted to 15°C

for one day until germination of the seed. Tissue is collected once the seedlings are 1 day old. At this point, the coleorhiza has just pushed out of the seed coat and the primary root is just making its appearance. The coleoptile has not yet pushed completely through the seed coat and is also just making its appearance. These 1 day old cold stressed seedlings are frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON~001 (Lib36, Lib83, Lib84) cDNA library is generated from maize leaves at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V8 stage. The older more juvenile leaves in a basal position as well as the younger more adult leaves which are more apical are all cut at the base, pooled and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMONN01 cDNA library is generated from maize (B73, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) normalized immature tassels at the V6 plant development stage normalized tissue. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into

10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V6 stage. At that stage the tassel is an immature tassel of about 2-3 cm in length. The tassels are removed and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN04 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) normalized total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10

inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN05 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) normalized root tissue at the V6 development

stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen and the harvested tissue is then stored at -80°C until RNA preparation. The single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN06 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) normalized total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-

hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The CMZ029 (SATMON036) cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) endosperm 22 days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of the maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergent to withhold the pollen. The ear shoots are pollinated and 22 days after pollination, the ears are pulled out and then the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the alurone layer is removed. After dissection, the endosperms are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The CMz030 (Lib143) cDNA library is generated from maize seedling tissue two days post germination. Seeds are planted on a moist filter paper on a covered try that is keep in the dark until germination. The trays are then moved to the bench top at 15 hr daytime/9 hr nighttime cycles for 2 days post-germination. The day time temperature is 80°F and the

nighttime temperature is 70°F. Tissue is collected when the seedlings are 2 days old. At this stage, the colehrhiza has pushed through the seed coat and the primary root (the radicle) is just piercing the colehrhiza and is barely visible. The seedlings are placed at 42°C for 1 hour. Following the heat shock treatment, the seedlings are immersed in liquid nitrogen and crushed. The harvested tissue is stored at -80° until RNA preparation.

The CMz031 (Lib148) cDNA library is generated from maize pollen tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag to withhold pollen. Twenty-one days after pollination, prior to removing the ears, the paper bag is shaken to collect the mature pollen. The mature pollen is immediately frozen in liquid nitrogen containers and the pollen is crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz033 (Lib189) cDNA library is generated from maize pooled leaf tissue. Samples are harvested from open pollinated plants. Tissue is collected from maize leaves at the

anthesis stage. The leaves are collected from 10-12 plants and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz034 (Lib3060) cDNA library is generated from maize mature tissue at 40 days post pollination plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from leaves located two leaves below the ear leaf. This sample represents those genes expressed during onset and early stages of leaf senescence. The leaves are pooled and immediately transferred to liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz036 (Lib3062) cDNA library is generated from maize husk tissue at the 8 week old plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during

the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from 8 week old plants. The husk is separated from the ear and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz037 (Lib3059) cDNA library is generated from maize pooled kernal at 12-15 days after pollination plant development stage. Sample were collected from field grown material. Whole kernal from hand pollinated (control pollination) are harvested as whole ears and immediately frozen on dry ice. Kernels from 10-12 ears were pooled and ground together in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz039 (Lib3066) cDNA library is generated from maize immature anther tissue at the 7 week old immature tassel stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 7 week old immature tassel stage. At this stage, prior to anthesis, the

immature anthers are green and enclosed in the staminate spikelet. The developing anthers are dissected away from the 7 week old immature tassel and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz040 (Lib3067) cDNA library is generated from maize kernel tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag before silk emergence to withhold pollen. Five to eight days after controlled pollination. The ears are pulled and the kernels removed. The kernels are immediately frozen in liquid nitrogen. The harvested kernels tissue is then stored at -80°C until RNA preparation. This sample represents gene expressed in early kernel development, during periods of cell division, amyloplast biogenesis and early carbon flow across the material to filial tissue.

The CMz041 (Lib3068) cDNA library is generated from maize pollen germinating silk tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are

transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants when the ear shoots are ready for fertilization at the silk emergence stage. The emerging silks are pollinated with an excess of pollen under controlled pollination conditions in the green house. Eighteen hours after pollination the silks are removed from the ears and immediately frozen in liquid nitrogen containers. This sample represents genes expressed in both pollen and silk tissue early in pollination. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz042 (Lib3069) cDNA library is generated from maize ear tissue excessively pollinated at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants and the ear shoots which are ready for fertilization are at the silk emergence stage. The immature ears are pollinated with an excess of pollen under controlled pollination conditions. Eighteen hours post-pollination, the ears are removed and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz044 (Lib3075) cDNA library is generated from maize microspore tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from immature anthers from 7 week old tassels. The immature anthers are first dissected from the 7 week old tassel with a scalpel on a glass slide covered with water. The microspores (immature pollen) are released into the water and are recovered by centrifugation. The microspore suspension is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz045 (Lib3076) cDNA library is generated from maize immature ear megaspore tissue. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing

Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from immature ear (megaspore) obtained from 7 week old plants. The immature ears are harvested from the 7 week old plants and are approximately 2.5 to 3 cm in length. The kernels are removed from the cob immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz047 (Lib3078) cDNA library is generated from maize CO₂ treated high-exposure shoot tissue at the V10+ plant development stage. RX601 maize seeds are sterilized for 1 minute with a 10% clorox solution. The seeds are rolled in germination paper, and germinated in 0.5 mM calcium sulfate solution for two days at 30°C. The seedlings are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium at a rate of 2-3 seedlings per pot. Twenty pots are placed into a high CO₂ environment (approximately 1000 ppm CO₂). Twenty plants were grown under ambient greenhouse CO₂ (approximately 450 ppm CO₂). Plants are watered daily before transplantation and three times a week after transplantation. Peters 20-20-20 fertilizer is also lightly applied. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime

temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. At ten days post planting, the shoots from both atmosphere are frozen in liquid nitrogen and lightly ground. The roots are washed in deionized water to remove the support media and the tissue is immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz048 (Lib3079) cDNA library is generated from maize basal endosperm transfer layer tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ maize plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag prior to silk emergence, to withhold the pollen. Kernels are harvested at 12 days post-pollination and placed on wet ice for dissection. The kernels are cross sectioned laterally, dissecting just above the pedicel region, including 1-2 mm of the lower endosperm and the basal endosperm transfer region. The pedicel and lower endosperm region containing the basal endosperm transfer layer is pooled and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is beyond the 10-leaf development stage and the ear shoots are approximately 15-20 cm in length. The ears are pulled and silks are separated from the ears and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON001 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) total leaf tissue at the V4 plant development stage. Leaf tissue from 38, field grown V4 stage plants is harvested from the 4th node. Leaf tissue is removed from the plants and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON002 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue at the V4 plant development stage. Root tissue from 76, field grown V4 stage plants is harvested. The root systems is cut from the soybean plant and washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON003 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling hypocotyl axis tissue harvested 2 day post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even

moisture conditions. Tissue is collected 2 days after the start of imbibition. The 2 days after imbibition samples are separated into 3 collections after removal of any adhering seed coat. At the 2 day stage, the hypocotyl axis is emerging from the soil. A few seedlings have cracked the soil surface and exhibited slight greening of the exposed cotyledons. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON004 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling cotyledon tissue harvested 2 day post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 2 days after the start of imbibition. The 2 days after imbibition samples are separated into 3 collections after removal of any adhering seed coat. At the 2 day stage, the hypocotyl axis is emerging from the soil. A few seedlings have cracked the soil surface and exhibited slight greening of the exposed cotyledons. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON005 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling hypocotyl axis tissue harvested 6 hour post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the

nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 6 hours after the start of imbibition. The 6 hours after imbibition samples are separated into 3 collections after removal of any adhering seed coat. The 6 hours after imbibition sample is collected over the course of approximately 2 hours starting at 6 hours post imbibition. At the 6 hours after imbibition stage, not all cotyledons have become fully hydrated and germination, or radicle protrusion, has not occurred. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON006 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling cotyledons tissue harvest 6 hour post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 6 hours after imbibition. The 6 hours after imbibition samples are separated into 3 collections after removal of any adhering seed coat. The 6 hours after imbibition sample is collected over the course of approximately 2 hours starting at 6 hours post-imbibition. At the 6 hours after imbibition, not all cotyledons have become fully hydrated and germination or radicle protrusion, have not occurred. The seedlings are washed in water to remove soil, cotyledon harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON007 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 and 35 days post-

flowering. Seed pods from field grown plants are harvested 25 and 35 days after flowering and the seeds extracted from the pods. Approximately 4.4g and 19.3g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON008 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested from 25 and 35 days post-flowering plants. Total leaf tissue is harvested from field grown plants. Approximately 19g and 29g of leaves are harvested from the fourth node of the plant 25 and 35 days post-flowering and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON009 cDNA library is generated from soybean cutlivar C1944 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) pod and seed tissue harvested 15 days post-flowering. Pods from field grown plants are harvested 15 days post-flowering. Approximately 3g of pod tissue is harvested and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON010 cDNA library is generated from soybean cultivar C1944 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) seed tissue harvested 40 days post-flowering. Pods from field grown plants are harvested 40 days post-flowering. Pods and seeds are separated, approximately 19g of seed tissue is harvested and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON011 cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical

germ plasma) leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4th node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON012 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue. Leaves from field grown plants are harvested from the fourth node 15 days post-flowering. Approximately 12g of leaves are harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON013 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root and nodule tissue. Approximately, 28g of root tissue from field grown plants is harvested 15 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON014 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 and 35 days after flowering. Seed pods from field grown plants are harvested 15 days after flowering and the seeds extracted from the pods. Approximately 5g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON015 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 45 and 55 days post-flowering. Seed pods from field grown plants are harvested 45 and 55 days after flowering and the seeds extracted from the pods. Approximately 19g and 31g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON016 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue. Approximately, 61g and 38g of root tissue from field grown plants is harvested 25 and 35 days post-flowering is harvested. The root system is cut from the soybean plant and washed with water to free it from the soil. The tissue is placed in 14ml polystyrene tubes and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON017 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue. Approximately 28g of root tissue from field grown plants is harvested 45 and 55 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON018 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested 45 and 55 days post-flowering. Leaves from field grown plants are harvested 45 and 55 days after flowering from the fourth node. Approximately 27g and 33g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON019 cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) root tissue. Roots are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 50g and 56g of roots are harvested from each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON020 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 65 and 75 days post-flowering. Seed pods from field grown plants are harvested 45 and 55 days after flowering and the seeds extracted from the pods. Approximately 14g and 31g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON021 cDNA library is generated from Soybean Cyst Nematode-resistant soybean cultivar Hartwig (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) root tissue. Plants are grown in tissue culture at room temperature. At approximately 6 weeks post-germination, the plants are exposed to sterilized Soybean Cyst Nematode eggs. Infection is then allowed to progress for 10 days. After the 10 day infection process, the tissue is harvested. Agar from the culture medium and nematodes are removed and the root tissue is immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON022 (Lib3030) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) partially opened flower tissue.

Partially to fully opened flower tissue is harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. A total of 3g of flower tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON023 cDNA library is generated from soybean genotype BW211S Null (Tohoku University, Morioka, Japan) seed tissue harvested 15 and 40 days post-flowering. Seed pods from field grown plants are harvested 15 and 40 days post-flowering and the seeds extracted from the pods. Approximately 0.7g and 14.2g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON024 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) internode-2 tissue harvested 18 days post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. The plants are grown in a greenhouse for 18 days after the start of imbibition at ambient temperature. Soil is checked and watered daily to maintain even moisture conditions. Stem tissue is harvested 18 days after the start of imbibition. The samples are divided into hypocotyl and internodes 1 through 5. The fifth internode contains some leaf bud material. Approximately 3 g of each sample is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON025 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested 65 days post-flowering. Leaves are harvested from the fourth node of field grown plants 65 days post-flowering.

Approximately 18.4g of leaf tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

SOYMON026 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue harvested 65 and 75 days post-flowering. Approximately 27g and 40g of root tissue from field grown plants is harvested 65 and 75 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON027 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 days post-flowering. Seed pods from field grown plants are harvested 25 days post-flowering and the seeds extracted from the pods. Approximately 17g of seeds are harvested from the seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON028 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought-stressed root tissue. The plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of development, water is withheld from half of the plant collection (drought stressed population). After 3 days, half of the plants from the drought stressed condition and half of the plants from the control population are harvested. After another 3 days (6 days post drought induction) the remaining plants are

harvested. A total of 27g and 40g of root tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON029 cDNA library is generated from Soybean Cyst Nematode-resistant soybean cultivar PI07354 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) root tissue. Late fall to early winter greenhouse grown plants are exposed to Soybean Cyst Nematode eggs. At 10 days post-infection, the plants are uprooted, rinsed briefly and the roots frozen in liquid nitrogen. Approximately 20 grams of root tissue is harvested from the infected plants. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON030 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) flower bud tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Flower buds are removed from the plant at the pedicel. A total of 100mg of flower buds are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON031 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) carpel and stamen tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Flower buds are

removed from the plant at the pedicel. Flowers are dissected to separate petals, sepals and reproductive structures (carpels and stamens). A total of 300mg of carpel and stamen tissue are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON032 cDNA library is prepared from the Asgrow cultivar A4922 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) rehydrated dry soybean seed meristem tissue. Surface sterilized seeds are germinated in liquid media for 24 hours. The seed axis is then excised from the barely germinating seed, placed on tissue culture media and incubated overnight at 20°C in the dark. The supportive tissue is removed from the explant prior to harvest. Approximately 570mg of tissue is harvested and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON033 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) heat-shocked seedling tissue without cotyledons. Seeds are imbibed and germinated in vermiculite for 2 days under constant illumination. After 48 hours, the seedlings are transferred to an incubator set at 40°C under constant illumination. After 30, 60 and 180 minutes seedlings are harvested and dissected. A portion of the seedling consisting of the root, hypocotyl and apical hook is frozen in liquid nitrogen and stored at -80°C. The seedlings after 2 days of imbibition are beginning to emerge from the vermiculite surface. The apical hooks are dark green in appearance. Total RNA and poly A⁺ RNA is prepared from equal amounts of pooled tissue.

The SOYMON034 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) cold-shocked seedling tissue without cotyledons. Seeds are imbibed and germinated in vermiculite for 2 days under constant

illumination. After 48 hours, the seedlings are transferred to a cold room set at 5°C under constant illumination. After 30, 60 and 180 minutes seedlings are harvested and dissected. A portion of the seedling consisting of the root, hypocotyl and apical hook is frozen in liquid nitrogen and stored at -80°C. The seedlings after 2 days of imbibition are beginning to emerge from the vermiculite surface. The apical hooks are dark green in appearance.

The SOYMON035 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed coat tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are harvested from mid to nearly full maturation (seed coats are not yellowing). The entire embryo proper is removed from the seed coat sample and the seed coat tissue are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON036 cDNA library is generated from soybean cultivars PI171451, PI227687 and PI229358 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) insect challenged leaves. Plants from each of the three cultivars are grown in screenhouse conditions. The screenhouse is divided in half and one half of the screenhouse is infested with soybean looper and the other half infested with velvetbean caterpillar. A single leaf is taken from each of the representative plants at 3 different time points, 11 days after infestation, 2 weeks after infestation and 5 weeks after infestation and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation. Total RNA and poly A+ RNA is

isolated from pooled tissue consisting of equal quantities of all 18 samples (3 genotypes X 3 sample times X 2 insect genotypes).

The SOYMON037 cDNA library is generated from soybean cultivar A3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) etiolated axis and radical tissue. Seeds are planted in moist vermiculite, wrapped and kept at room temperature in complete darkness until harvest. Etiolated axis and hypocotyl tissue is harvested at 2, 3 and 4 days post-planting. A total of 1 gram of each tissue type is harvested at 2, 3 and 4 days after planting and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON038 cDNA library is generated from soybean variety Asgrow A3237 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) rehydrated dry seeds. Explants are prepared for transformation after germination of surface-sterilized seeds on solid tissue media. After 6 days, at 28°C and 18 hours of light per day, the germinated seeds are cold shocked at 4°C for 24 hours. Meristemic tissue and part of the hypocotyl is removed and cotyledon excised. The prepared explant is then wounded for *Agrobacterium* infection. The 2 grams of harvested tissue is frozen in liquid nitrogen and stored at -80°C until RNA preparation.

The Soy51 (LIB3027) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The

dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The Soy52 (LIB3028) cDNA library is generated from normalized flower DNA. Single stranded DNA representing approximately 1×10^6 colony forming units of SOYMON022 harvested tissue is used as the starting material for normalization. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The Soy53 (LIB3039) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling shoot apical meristem tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Apical tissue is harvested from seedling shoot meristem tissue, 7-8 days after the start of imbibition. The apex of each seedling is dissected to include the fifth node to the apical meristem. The fifth node corresponds to the third trifoliate leaf in the very early stages of development. Stipules completely envelop the leaf primordia at this time. A total of 200mg of apical tissue is harvested

and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The Soy54 (LIB3040) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) heart to torpedo stage embryo tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are collected and embryos removed from surrounding endosperm and maternal tissues. Embryos from globular to young torpedo stages (by corresponding analogy to *Arabidopsis*) are collected with a bias towards the middle of this spectrum. Embryos which are beginning to show asymmetric development of cotyledons are considered the upper developmental boundary for the collection and are excluded. A total of 12 mg embryo tissue is frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy55 (LIB3049) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) young seed tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are collected from very young pods (5 to 15 days after flowering). A total of 100mg of seeds are harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy56 (LIB3029) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are not converted to double stranded form and represent a non-normalized seed pool for comparison to Soy51 cDNA libraries.

The Soy58 (LIB3050) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed root tissue subtracted from control root tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days root tissue from both drought stressed and control (watered regularly) plants are collected and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that

described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy59 (LIB3051) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) endosperm tissue. Seeds are germinated on paper towels under laboratory ambient light conditions. At 8, 10 and 14 hours after imbibition, the seed coats are harvested. The endosperm consists of a very thin layer of tissue affixed to the inside of the seed coat. The seed coat and endosperm are frozen immediately after harvest in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The Soy60 (LIB3072) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed seed plus pod subtracted from control seed plus pod tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and

control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy61 (LIB3073) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. After 18hours, 24hours and 48 hours post

treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). For this library's construction, the eighth fraction of the cDNA size fractionation step was used for ligation.

The Soy62 (LIB3074) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St.

Loius, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. After 18 hours, 24 hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). For this library's construction, the ninth fraction of the cDNA size fractionation step was used for ligation.

The Soy65 (LIB3107) 07cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought-stressed abscission zone tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr

nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Plants are irrigated with 15-16-17 Peter's Mix. At the R3 stage of development, drought is imposed by withholding water. At 3, 4, 5 and 6 days, tissue is harvested and wilting is not obvious until the fourth day. Abscission layers from reproductive organs are harvested by cutting less than one millimeter proximal and distal to the layer and immediately frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The Soy66 (LIB3109) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) non-drought stressed abscission zone tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Plants are irrigated with 15-16-17 Peter's Mix. At 3, 4, 5 and 6 days, control abscission layer tissue is harvested. Abscission layers from reproductive organs are harvested by cutting less than one millimeter proximal and distal to the layer and immediately frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy67 (LIB3065) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar

ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. Captured hybrids are eluted with water.

Soy68 (LIB3052) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. Captured hybrids are eluted with water.

Soy69 (LIB3053) cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) normalized leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4th node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately 1×10^6 colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the

synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

Soy70 (LIB3055) cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4th node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

Soy71 (LIB3056) cDNA library is generated from soybean cultivars Cristalina and FT108 (tropical germ plasma) root tissue. Roots are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 50g and 56g of roots are harvested from each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

Soy72 (LIB3093) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed leaf control tissue. Seeds

are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

Soy73 (LIB3093) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed leaf subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under

12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy76 (Lib3106) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid and arachidonic treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the

plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. Arachidonic treated seedlings are sprayed with 1m/ml arachidonic acid in 0.1% Tween-20. After 18hours, 24hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. The RNA from the arachidonic treated seedlings is isolated separately. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). Fraction 10 of the size fractionated cDNA is ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.) in order to capture some of the smaller transcripts characteristic of antifungal proteins.

Soy77 (LIB3108) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. Arachidonic treated seedlings are sprayed with 1m/ml arachidonic acid in 0.1% Tween-20. After 18 hours, 24 hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. The RNA from the arachidonic treated seedlings is isolated separately. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After

hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). Fraction 10 of the size fractionated cDNA is ligated into the pSPORT vector in order to capture some of the smaller transcripts characteristic of antifungal proteins.

The stored RNA is purified using Trizol reagent from Life Technologies (Gibco BRL, Life Technologies, Gaithersburg, Maryland U.S.A.), essentially as recommended by the manufacturer. Poly A⁺ RNA (mRNA) is purified using magnetic oligo dT beads essentially as recommended by the manufacturer (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.).

Construction of plant cDNA libraries is well-known in the art and a number of cloning strategies exist. A number of cDNA library construction kits are commercially available. The SuperscriptTM Plasmid System for cDNA synthesis and Plasmid Cloning (Gibco BRL, Life Technologies, Gaithersburg, Maryland U.S.A.) is used, following the conditions suggested by the manufacturer.

Normalized libraries are made using essentially the Soares procedure (Soares *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:9228-9232 (1994), the entirety of which is herein incorporated by reference). This approach is designed to reduce the initial 10,000-fold variation in individual cDNA frequencies to achieve abundances within one order of magnitude while maintaining the overall sequence complexity of the library. In the normalization process, the prevalence of high-abundance cDNA clones decreases dramatically, clones with mid-level abundance are relatively unaffected and clones for rare transcripts are effectively increased in abundance.

Example 2

The cDNA libraries are plated on LB agar containing the appropriate antibiotics for selection and incubated at 37° for a sufficient time to allow the growth of individual colonies. Single colonies are individually placed in each well of a 96-well microtiter plates containing LB liquid including the selective antibiotics. The plates are incubated overnight at approximately 37°C with gentle shaking to promote growth of the cultures. The plasmid DNA is isolated from each clone using Qiaprep plasmid isolation kits, using the conditions recommended by the manufacturer (Qiagen Inc., Santa Clara, California U.S.A.).

Template plasmid DNA clones are used for subsequent sequencing. For sequencing, the ABI PRISM dRhodamine Terminator Cycle Sequencing Ready Reaction Kit with AmpliTaq® DNA Polymerase, FS, is used (PE Applied Biosystems, Foster City, California U.S.A.).

Example 3

Nucleic acid sequences that encode for the following proteins: methionine adenosyltransferase, S-adenosylmethionine decarboxylase, aspartate kinase, aspartate-semialdehyde dehydrogenase, *O*-succinylhomoserine (thiol)-lyase, cystathionine β -lyase, 5-methyltetrahydropteroyltriglutamate, adenosylhomocysteinase, cystathionine β -synthase, cystathionine γ -lyase and *O*-acetylhomoserine (thiol)-lyase are identified from the Monsanto EST PhytoSeq database using TBLASTN (default values)(TBLASTN compares a protein query against the six reading frames of a nucleic acid sequence). Matches found with BLAST P values equal or less than 0.001 (probability) or BLAST Score of equal or greater than 90 are classified as hits. If the program used to determine the hit is HMMSW then the score refers to HMMSW score.

In addition, the GenBank database is searched with BLASTN and BLASTX (default values) using ESTs as queries. EST that pass the hit probability threshold of $10e^{-8}$ for the following enzymes are combined with the hits generated by using TBLASTN (described above) and classified by enzyme (see Table A below).

A cluster refers to a set of overlapping clones in the PhytoSeq database. Such an overlapping relationship among clones is designated as a “cluster” when BLAST scores from pairwise sequence comparisons of the member clones meets a predetermined minimum value or product score of 50 or more (Product Score = (BLAST SCORE x Percentage Identity)/(5 x minimum [length (Seq1), length (Seq2)]))

Since clusters are formed on the basis of single-linkage relationships, it is possible for two non-overlapping clones to be members of the same cluster if, for instance, they both overlap a third clone with at least the predetermined minimum BLAST score (stringency). A cluster ID is arbitrarily assigned to all of those clones which belong to the same cluster at a given stringency and a particular clone will belong to only one cluster at a given stringency. If a cluster contains only a single clone (a “singleton”), then the cluster ID number will be negative, with an absolute value equal to the clone ID number of its single member. Clones grouped in a cluster in most cases represent a contiguous sequence.

TABLE A*

METHIONINE ADENOSYLTRANSFERASE (EC 2.5.1.6)								
Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1	-700019427	700019427H1	SATMON001	g17262	BLASTX	120	1e-11	92
2	-700074004	700074004H1	SATMON007	g1778820	BLASTN	830	1e-60	80
3	-700149718	700149718H1	SATMON007	g1778820	BLASTN	263	1e-13	80
4	-700220251	700220251H1	SATMON011	g1127582	BLASTN	243	1e-9	90
5	-700458196	700458196H1	SATMON029	g1778820	BLASTN	202	1e-22	88
6	-701172459	701172459H1	SATMONN05	g882334	BLASTN	801	1e-57	89
7	-L1436317	LIB143-062-Q1-E1-B6	LIB143	g1778820	BLASTN	341	1e-19	77
8	-L1893126	LIB189-021-Q1-E1-F4	LIB189	g1778820	BLASTN	629	1e-41	88
9	-L1894542	LIB189-033-Q1-E1-H12	LIB189	g1778820	BLASTN	564	1e-60	76
10	-L30622839	LIB3062-028-Q1-K1-C2	LIB3062	g1778820	BLASTN	625	1e-65	74
11	-L30671966	LIB3067-036-Q1-K1-D8	LIB3067	g1655576	BLASTX	118	1e-25	85
12	-L30682166	LIB3068-004-Q1-K1-D5	LIB3068	g497900	BLASTX	111	1e-29	47
13	1	LIB143-036-Q1-E1-G7	LIB143	g450548	BLASTN	1766	1e-138	88
14	1	LIB3060-013-Q1-K1-F7	LIB3060	g1778820	BLASTN	1702	1e-133	90
15	1	LIB143-013-Q1-E1-A6	LIB143	g1778820	BLASTN	1712	1e-133	88
16	1	LIB148-007-Q1-E1-B8	LIB148	g450548	BLASTN	1693	1e-132	88
17	1	LIB3079-003-Q1-K1-G1	LIB3079	g1778820	BLASTN	1236	1e-130	88
18	1	LIB3066-011-Q1-K1-E4	LIB3066	g1778820	BLASTN	1661	1e-129	86
19	1	LIB3068-008-Q1-K1-D2	LIB3068	g450548	BLASTN	1192	1e-128	88
20	1	LIB143-061-Q1-E1-E7	LIB143	g1778820	BLASTN	1638	1e-127	88
21	1	LIB189-006-Q1-E1-B2	LIB189	g1778820	BLASTN	1627	1e-126	86
22	1	LIB143-031-Q1-E1-C7	LIB143	g450548	BLASTN	1595	1e-124	86
23	1	LIB3068-011-Q1-K1-B12	LIB3068	g450548	BLASTN	1552	1e-123	87
24	1	LIB143-013-Q1-E1-G11	LIB143	g450548	BLASTN	1585	1e-123	88
25	1	LIB3062-057-Q1-K1-H9	LIB3062	g450548	BLASTN	1577	1e-122	89
26	1	LIB143-008-Q1-E1-G9	LIB143	g450548	BLASTN	1556	1e-120	89
27	1	LIB3068-058-Q1-K1-B7	LIB3068	g450548	BLASTN	1470	1e-118	86
28	1	LIB3067-048-Q1-K1-G5	LIB3067	g1778820	BLASTN	1518	1e-117	89
29	1	LIB143-049-Q1-E1-A12	LIB143	g1778820	BLASTN	1480	1e-114	91
30	1	LIB3062-010-Q1-K1-C7	LIB3062	g1778820	BLASTN	1145	1e-113	87
31	1	LIB3068-055-Q1-K1-E10	LIB3068	g450548	BLASTN	1451	1e-112	88
32	1	LIB189-025-Q1-E1-H1	LIB189	g1778820	BLASTN	1460	1e-112	90
33	1	LIB143-066-Q1-E1-F4	LIB143	g1778820	BLASTN	1309	1e-111	86
34	1	LIB143-003-Q1-E1-D10	LIB143	g1778820	BLASTN	1440	1e-111	87
35	1	LIB3062-033-Q1-K1-G1	LIB3062	g1778820	BLASTN	1449	1e-111	86
36	1	LIB3066-029-Q1-K1-H11	LIB3066	g1778820	BLASTN	1427	1e-110	88
37	1	LIB3068-016-Q1-K1-D9	LIB3068	g450548	BLASTN	1213	1e-109	87
38	1	LIB3062-027-Q1-K1-B11	LIB3062	g1778820	BLASTN	1187	1e-108	87
39	1	LIB3068-048-Q1-K1-G4	LIB3068	g1778820	BLASTN	1410	1e-108	82
40	1	LIB148-020-Q1-E1-B6	LIB148	g1778820	BLASTN	1088	1e-107	84
41	1	LIB3066-046-Q1-K1-F2	LIB3066	g1778820	BLASTN	1400	1e-107	90
42	1	700084130H1	SATMON011	g1778820	BLASTN	960	1e-106	92
43	1	700092863H1	SATMON008	g1778820	BLASTN	1388	1e-106	92
44	1	LIB3068-043-Q1-K1-A12	LIB3068	g450548	BLASTN	827	1e-105	81
45	1	700092659H1	SATMON008	g1778820	BLASTN	1359	1e-104	92
46	1	LIB143-038-Q1-E1-H11	LIB143	g450548	BLASTN	1345	1e-103	84
47	1	LIB3062-028-Q1-K1-F12	LIB3062	g1778820	BLASTN	1082	1e-102	82
48	1	700103135H1	SATMON010	g1778820	BLASTN	1306	1e-100	93

49	1	LIB3078-050-Q1-K1-D9	LIB3078	g1778820	BLASTN	1309	1e-100	86
50	1	700084823H1	SATMON011	g1778820	BLASTN	1311	1e-100	90
51	1	700201311H1	SATMON003	g1778820	BLASTN	960	1e-97	89
52	1	700265914H1	SATMON017	g1778820	BLASTN	1275	1e-97	91
53	1	700205948H1	SATMON003	g1778820	BLASTN	983	1e-96	93
54	1	LIB189-003-Q1-E1-A6	LIB189	g450548	BLASTN	1262	1e-96	86
55	1	LIB143-017-Q1-E1-B9	LIB143	g1778820	BLASTN	779	1e-95	86
56	1	700097434H1	SATMON009	g450548	BLASTN	1247	1e-95	90
57	1	700089089H1	SATMON011	g450548	BLASTN	1251	1e-95	89
58	1	700071949H1	SATMON007	g1778820	BLASTN	1256	1e-95	87
59	1	700047892H1	SATMON003	g1778820	BLASTN	1236	1e-94	91
60	1	700077246H1	SATMON007	g450548	BLASTN	1237	1e-94	89
61	1	700085681H1	SATMON011	g1778820	BLASTN	1237	1e-94	89
62	1	LIB3067-010-Q1-K1-A7	LIB3067	g1778820	BLASTN	1244	1e-94	83
63	1	700619961H1	SATMON034	g450548	BLASTN	962	1e-93	89
64	1	700087395H1	SATMON011	g450548	BLASTN	1040	1e-93	89
65	1	700240431H1	SATMON010	g1778820	BLASTN	1156	1e-93	92
66	1	700053822H1	SATMON011	g1778820	BLASTN	1224	1e-93	91
67	1	700103856H1	SATMON010	g450548	BLASTN	1224	1e-93	88
68	1	700083703H1	SATMON011	g450548	BLASTN	1229	1e-93	89
69	1	700087360H1	SATMON011	g450548	BLASTN	1234	1e-93	90
70	1	700104713H1	SATMON010	g450548	BLASTN	1188	1e-92	87
71	1	700265996H1	SATMON017	g450548	BLASTN	1214	1e-92	88
72	1	700264870H1	SATMON017	g450548	BLASTN	945	1e-91	89
73	1	700219348H1	SATMON011	g1778820	BLASTN	1203	1e-91	91
74	1	700095527H1	SATMON008	g450548	BLASTN	1153	1e-90	88
75	1	LIB3068-023-Q1-K1-G5	LIB3068	g1778820	BLASTN	1187	1e-90	89
76	1	700102547H1	SATMON010	g1778820	BLASTN	1194	1e-90	88
77	1	700332179H1	SATMON019	g450548	BLASTN	1054	1e-89	88
78	1	700405470H1	SATMON029	g450548	BLASTN	1071	1e-89	90
79	1	LIB3069-043-Q1-K1-G2	LIB3069	g450548	BLASTN	1175	1e-89	84
80	1	700451332H1	SATMON028	g450548	BLASTN	1176	1e-89	90
81	1	700028108H1	SATMON003	g450548	BLASTN	1180	1e-89	89
82	1	700076154H1	SATMON007	g1778820	BLASTN	1180	1e-89	89
83	1	700089770H1	SATMON011	g450548	BLASTN	1185	1e-89	89
84	1	700051882H1	SATMON003	g1778820	BLASTN	844	1e-88	91
85	1	700242942H1	SATMON010	g1778820	BLASTN	1163	1e-88	91
86	1	700094996H1	SATMON008	g1778820	BLASTN	1166	1e-88	88
87	1	700476239H1	SATMON025	g450548	BLASTN	1166	1e-88	89
88	1	700219530H1	SATMON011	g1778820	BLASTN	1005	1e-87	91
89	1	700049726H1	SATMON003	g1778820	BLASTN	1041	1e-87	89
90	1	700096904H1	SATMON008	g1778820	BLASTN	1157	1e-87	86
91	1	700344026H1	SATMON021	g450548	BLASTN	1158	1e-87	88
92	1	700105590H1	SATMON010	g1778820	BLASTN	1139	1e-86	89
93	1	700465283H1	SATMON025	g450548	BLASTN	1145	1e-86	89
94	1	700104341H1	SATMON010	g450548	BLASTN	1146	1e-86	87
95	1	LIB3062-021-Q1-K1-F12	LIB3062	g1778820	BLASTN	923	1e-85	78
96	1	700105246H1	SATMON010	g450548	BLASTN	930	1e-85	89
97	1	700072146H1	SATMON007	g1778820	BLASTN	1131	1e-85	91
98	1	700028553H1	SATMON003	g1778820	BLASTN	1132	1e-85	88
99	1	700451419H1	SATMON028	g450548	BLASTN	1133	1e-85	89
100	1	700103241H1	SATMON010	g450548	BLASTN	1134	1e-85	90
101	1	700612549H1	SATMON033	g450548	BLASTN	1134	1e-85	89
102	1	700050434H1	SATMON003	g450548	BLASTN	723	1e-84	89

103	1	700094764H1	SATMON008	g1778820	BLASTN	977	1e-84	86
104	1	700163030H1	SATMON013	g1778820	BLASTN	1011	1e-84	93
105	1	700466542H1	SATMON025	g450548	BLASTN	1085	1e-84	89
106	1	700047380H1	SATMON003	g450548	BLASTN	1096	1e-84	89
107	1	700456429H1	SATMON029	g450548	BLASTN	1118	1e-84	88
108	1	700096049H1	SATMON008	g1778820	BLASTN	1122	1e-84	86
109	1	700075906H1	SATMON007	g1778820	BLASTN	838	1e-83	86
110	1	LIB3067-048-Q1-K1-G3	LIB3067	g1778820	BLASTN	1045	1e-83	86
111	1	700214043H1	SATMON016	g450548	BLASTN	1103	1e-83	88
112	1	700096909H1	SATMON008	g1778820	BLASTN	1103	1e-83	90
113	1	701184635H1	SATMONN06	g1778820	BLASTN	1104	1e-83	87
114	1	700158969H1	SATMON012	g1778820	BLASTN	1107	1e-83	93
115	1	700475902H1	SATMON025	g450548	BLASTN	1112	1e-83	89
116	1	700207076H1	SATMON003	g1778820	BLASTN	1113	1e-83	90
117	1	700161901H1	SATMON012	g1778820	BLASTN	1091	1e-82	91
118	1	700452183H1	SATMON028	g450548	BLASTN	1095	1e-82	88
119	1	700452326H1	SATMON028	g450548	BLASTN	1098	1e-82	88
120	1	700106918H1	SATMON010	g450548	BLASTN	1101	1e-82	88
121	1	700220146H1	SATMON011	g450548	BLASTN	1083	1e-81	89
122	1	700221060H1	SATMON011	g450548	BLASTN	1084	1e-81	88
123	1	700243344H1	SATMON010	g1778820	BLASTN	1084	1e-81	92
124	1	700475790H1	SATMON025	g450548	BLASTN	1084	1e-81	89
125	1	700452691H1	SATMON028	g1778820	BLASTN	1086	1e-81	87
126	1	700243733H1	SATMON010	g450548	BLASTN	1086	1e-81	90
127	1	LIB3059-018-Q1-K1-D10	LIB3059	g1778820	BLASTN	830	1e-80	91
128	1	700165958H1	SATMON013	g450548	BLASTN	1070	1e-80	89
129	1	700082659H1	SATMON011	g1778820	BLASTN	1071	1e-80	88
130	1	700172557H1	SATMON013	g1778820	BLASTN	1074	1e-80	92
131	1	700221526H1	SATMON011	g450548	BLASTN	1075	1e-80	88
132	1	700456696H1	SATMON029	g450548	BLASTN	1077	1e-80	89
133	1	700430776H1	SATMONN01	g1778820	BLASTN	574	1e-79	90
134	1	LIB3062-025-Q1-K1-B11	LIB3062	g450548	BLASTN	585	1e-79	85
135	1	700077126H1	SATMON007	g1778820	BLASTN	743	1e-79	88
136	1	700160310H1	SATMON012	g1778820	BLASTN	752	1e-79	91
137	1	700030466H1	SATMON003	g1778820	BLASTN	757	1e-79	83
138	1	700210204H1	SATMON016	g1778820	BLASTN	844	1e-79	91
139	1	700094716H1	SATMON008	g1778820	BLASTN	857	1e-79	86
140	1	700157051H1	SATMON012	g1778820	BLASTN	937	1e-79	89
141	1	700241415H1	SATMON010	g450548	BLASTN	1055	1e-79	90
142	1	700160127H1	SATMON012	g450548	BLASTN	1056	1e-79	88
143	1	700205485H1	SATMON003	g1778820	BLASTN	1059	1e-79	90
144	1	700475930H1	SATMON025	g450548	BLASTN	1060	1e-79	89
145	1	700161902H1	SATMON012	g1778820	BLASTN	1060	1e-79	91
146	1	700213573H1	SATMON016	g450548	BLASTN	1061	1e-79	89
147	1	700207677H1	SATMON016	g450548	BLASTN	1061	1e-79	88
148	1	700581117H1	SATMON031	g450548	BLASTN	1065	1e-79	88
149	1	700451418H1	SATMON028	g450548	BLASTN	546	1e-78	89
150	1	700212606H1	SATMON016	g1778820	BLASTN	1044	1e-78	85
151	1	700222605H1	SATMON011	g1778820	BLASTN	1047	1e-78	86
152	1	700050542H1	SATMON003	g1778820	BLASTN	1053	1e-78	86
153	1	700450709H1	SATMON028	g450548	BLASTN	834	1e-77	88
154	1	700213368H1	SATMON016	g450548	BLASTN	861	1e-77	89
155	1	700095926H1	SATMON008	g1778820	BLASTN	1030	1e-77	86
156	1	700261686H1	SATMON017	g1778820	BLASTN	1034	1e-77	89

157	1	700235993H1	SATMON010	g450548	BLASTN	1037	1e-77	88
158	1	700466152H1	SATMON025	g450548	BLASTN	1038	1e-77	89
159	1	700266410H1	SATMON017	g1778820	BLASTN	1041	1e-77	85
160	1	700216991H1	SATMON016	g1778820	BLASTN	491	1e-76	92
161	1	700464957H1	SATMON025	g450548	BLASTN	607	1e-76	88
162	1	700160463H1	SATMON012	g1778820	BLASTN	1018	1e-76	91
163	1	700158224H1	SATMON012	g1778820	BLASTN	1018	1e-76	92
164	1	700469971H1	SATMON025	g1778820	BLASTN	1020	1e-76	86
165	1	700222731H1	SATMON011	g450548	BLASTN	1022	1e-76	88
166	1	700087434H1	SATMON011	g1778820	BLASTN	1023	1e-76	86
167	1	700094482H1	SATMON008	g1778820	BLASTN	1023	1e-76	86
168	1	700235976H1	SATMON010	g450548	BLASTN	1028	1e-76	89
169	1	700088266H1	SATMON011	g1778820	BLASTN	983	1e-75	92
170	1	700075882H1	SATMON007	g1778820	BLASTN	1009	1e-75	83
171	1	700243324H1	SATMON010	g1778820	BLASTN	1012	1e-75	86
172	1	700159625H2	SATMON012	g450548	BLASTN	1015	1e-75	90
173	1	700204422H1	SATMON003	g1778820	BLASTN	1015	1e-75	88
174	1	700048228H1	SATMON003	g1778820	BLASTN	741	1e-74	85
175	1	700072474H1	SATMON007	g1778820	BLASTN	799	1e-74	85
176	1	700477790H1	SATMON025	g450548	BLASTN	960	1e-74	89
177	1	LIB3059-022-Q1-K1-A9	LIB3059	g1778820	BLASTN	995	1e-74	86
178	1	700241633H1	SATMON010	g450548	BLASTN	997	1e-74	89
179	1	700093978H1	SATMON008	g1778820	BLASTN	999	1e-74	89
180	1	700238569H1	SATMON010	g450548	BLASTN	844	1e-73	88
181	1	700455160H1	SATMON029	g450548	BLASTN	849	1e-73	86
182	1	700262745H1	SATMON017	g1778820	BLASTN	984	1e-73	85
183	1	700405029H1	SATMON027	g1778820	BLASTN	990	1e-73	90
184	1	700195230H1	SATMON014	g450548	BLASTN	991	1e-73	88
185	1	700264753H1	SATMON017	g1778820	BLASTN	993	1e-73	90
186	1	700050142H1	SATMON003	g1778820	BLASTN	433	1e-72	86
187	1	700452677H1	SATMON028	g450548	BLASTN	919	1e-72	87
188	1	700168870H1	SATMON013	g1778820	BLASTN	971	1e-72	90
189	1	700023193H1	SATMON003	g450548	BLASTN	972	1e-72	88
190	1	700086582H1	SATMON011	g1778820	BLASTN	980	1e-72	84
191	1	700457710H1	SATMON029	g450548	BLASTN	982	1e-72	87
192	1	700477601H1	SATMON025	g450548	BLASTN	635	1e-71	88
193	1	700377730H1	SATMON019	g1778820	BLASTN	691	1e-71	87
194	1	700169782H1	SATMON013	g1778820	BLASTN	960	1e-71	91
195	1	700461113H1	SATMON033	g1778820	BLASTN	962	1e-71	85
196	1	700258669H1	SATMON017	g1778820	BLASTN	963	1e-71	90
197	1	700454253H1	SATMON029	g1778820	BLASTN	964	1e-71	85
198	1	700092376H1	SATMON008	g1778820	BLASTN	780	1e-70	83
199	1	700242885H1	SATMON010	g960356	BLASTN	946	1e-70	86
200	1	700094333H1	SATMON008	g1778820	BLASTN	947	1e-70	86
201	1	700575124H1	SATMON030	g450548	BLASTN	948	1e-70	82
202	1	700072544H1	SATMON007	g1778820	BLASTN	949	1e-70	86
203	1	LIB143-046-Q1-E1-B5	LIB143	g450548	BLASTN	951	1e-70	87
204	1	700096534H1	SATMON008	g1778820	BLASTN	956	1e-70	85
205	1	700262906H1	SATMON017	g1778820	BLASTN	937	1e-69	87
206	1	700094478H1	SATMON008	g1778820	BLASTN	939	1e-69	90
207	1	700172985H1	SATMON013	g450548	BLASTN	939	1e-69	88
208	1	700097938H1	SATMON009	g1778820	BLASTN	941	1e-69	85
209	1	LIB3069-006-Q1-K1-D5	LIB3069	g450548	BLASTN	943	1e-69	88
210	1	700093125H1	SATMON008	g1778820	BLASTN	945	1e-69	88

211	1	700424156H1	SATMONN01	g450548	BLASTN	752	1e-68	85
212	1	700405486H1	SATMON029	g450548	BLASTN	864	1e-68	91
213	1	LIB189-026-Q1-E1-H12	LIB189	g1778820	BLASTN	879	1e-68	87
214	1	700106172H1	SATMON010	g1778820	BLASTN	925	1e-68	85
215	1	700096890H1	SATMON008	g1778820	BLASTN	926	1e-68	90
216	1	700154404H1	SATMON007	g450548	BLASTN	928	1e-68	89
217	1	700468337H1	SATMON025	g450548	BLASTN	445	1e-67	88
218	1	700158827H1	SATMON012	g1778820	BLASTN	522	1e-67	87
219	1	700241744H1	SATMON010	g1778820	BLASTN	575	1e-67	85
220	1	700624633H1	SATMON034	g960356	BLASTN	644	1e-67	85
221	1	700154564H1	SATMON007	g1778820	BLASTN	735	1e-67	85
222	1	700172424H1	SATMON013	g450548	BLASTN	912	1e-67	90
223	1	700096287H1	SATMON008	g1778820	BLASTN	914	1e-67	86
224	1	700207638H1	SATMON016	g1778820	BLASTN	914	1e-67	86
225	1	700093735H1	SATMON008	g960356	BLASTN	920	1e-67	89
226	1	700159494H1	SATMON012	g1778820	BLASTN	899	1e-66	89
227	1	700236013H1	SATMON010	g1778820	BLASTN	900	1e-66	83
228	1	700220267H1	SATMON011	g1778820	BLASTN	439	1e-65	84
229	1	700477578H1	SATMON025	g450548	BLASTN	576	1e-65	90
230	1	700047743H1	SATMON003	g1778820	BLASTN	601	1e-65	82
231	1	700343041H1	SATMON021	g1778820	BLASTN	633	1e-65	86
232	1	700159886H1	SATMON012	g450548	BLASTN	803	1e-65	85
233	1	700570242H1	SATMON030	g450548	BLASTN	834	1e-65	85
234	1	700021930H1	SATMON001	g1778820	BLASTN	888	1e-65	87
235	1	700454959H1	SATMON029	g450548	BLASTN	890	1e-65	89
236	1	700171336H1	SATMON013	g1778820	BLASTN	890	1e-65	89
237	1	700105361H1	SATMON010	g1778820	BLASTN	806	1e-64	87
238	1	LIB3068-031-Q1-K1-B1	LIB3068	g450548	BLASTN	853	1e-64	89
239	1	700236324H1	SATMON010	g450548	BLASTN	875	1e-64	89
240	1	700150620H1	SATMON007	g450548	BLASTN	880	1e-64	87
241	1	700220648H1	SATMON011	g450548	BLASTN	881	1e-64	90
242	1	700150733H1	SATMON007	g450548	BLASTN	881	1e-64	85
243	1	700157367H1	SATMON012	g1778820	BLASTN	882	1e-64	85
244	1	700259676H1	SATMON017	g1778820	BLASTN	885	1e-64	88
245	1	700616490H1	SATMON033	g450548	BLASTN	886	1e-64	82
246	1	700102511H1	SATMON010	g450548	BLASTN	695	1e-63	89
247	1	700202805H1	SATMON003	g1778820	BLASTN	817	1e-63	92
248	1	700105685H1	SATMON010	g1778820	BLASTN	866	1e-63	84
249	1	700106113H1	SATMON010	g450548	BLASTN	873	1e-63	91
250	1	700444778H1	SATMON027	g1778820	BLASTN	343	1e-62	84
251	1	700103584H1	SATMON010	g450548	BLASTN	521	1e-62	86
252	1	LIB189-008-Q1-E1-D9	LIB189	g1778820	BLASTN	850	1e-62	91
253	1	700155684H1	SATMON007	g1778820	BLASTN	852	1e-62	85
254	1	700261639H1	SATMON017	g1778820	BLASTN	853	1e-62	89
255	1	700158367H1	SATMON012	g450548	BLASTN	856	1e-62	81
256	1	700153242H1	SATMON007	g1778820	BLASTN	859	1e-62	90
257	1	700210738H1	SATMON016	g1778820	BLASTN	859	1e-62	90
258	1	700203008H1	SATMON003	g450548	BLASTN	698	1e-61	86
259	1	700206081H1	SATMON003	g1778820	BLASTN	840	1e-61	92
260	1	700028643H1	SATMON003	g1778820	BLASTN	846	1e-61	85
261	1	700223914H1	SATMON011	g1778820	BLASTN	849	1e-61	84
262	1	700571455H1	SATMON030	g1778820	BLASTN	378	1e-60	88
263	1	700075374H1	SATMON007	g1778820	BLASTN	830	1e-60	85
264	1	700150452H1	SATMON007	g450548	BLASTN	831	1e-60	89

265	1	700261318H1	SATMON017	g1778820	BLASTN	831	1e-60	83
266	1	700616390H1	SATMON033	g450548	BLASTN	835	1e-60	92
267	1	700208718H1	SATMON016	g450548	BLASTN	561	1e-59	89
268	1	700448948H1	SATMON028	g1778820	BLASTN	653	1e-59	83
269	1	LIB3060-035-Q1-K1-H3	LIB3060	g450548	BLASTN	734	1e-59	85
270	1	700049753H1	SATMON003	g450548	BLASTN	769	1e-59	90
271	1	700154489H1	SATMON007	g1778820	BLASTN	814	1e-59	83
272	1	700170783H1	SATMON013	g1778820	BLASTN	816	1e-59	87
273	1	700237571H1	SATMON010	g450548	BLASTN	817	1e-59	89
274	1	700154872H1	SATMON007	g1778820	BLASTN	822	1e-59	85
275	1	700025620H1	SATMON004	g1778820	BLASTN	686	1e-58	84
276	1	700158255H1	SATMON012	g450548	BLASTN	803	1e-58	86
277	1	700159282H1	SATMON012	g450548	BLASTN	805	1e-58	83
278	1	700222171H1	SATMON011	g1778820	BLASTN	807	1e-58	89
279	1	700205003H1	SATMON003	g1778820	BLASTN	441	1e-57	86
280	1	700049209H1	SATMON003	g1778820	BLASTN	443	1e-57	90
281	1	700016430H1	SATMON001	g1778820	BLASTN	492	1e-57	86
282	1	700212936H1	SATMON016	g450548	BLASTN	564	1e-57	86
283	1	700156939H1	SATMON012	g1778820	BLASTN	801	1e-57	84
284	1	700222183H1	SATMON011	g1778820	BLASTN	561	1e-56	83
285	1	700203453H1	SATMON003	g1778820	BLASTN	784	1e-56	90
286	1	700167708H1	SATMON013	g1778820	BLASTN	560	1e-55	83
287	1	700214356H1	SATMON016	g1778820	BLASTN	693	1e-55	87
288	1	700475377H1	SATMON025	g450548	BLASTN	705	1e-55	90
289	1	700029625H1	SATMON003	g450548	BLASTN	712	1e-55	88
290	1	700202091H1	SATMON003	g1778820	BLASTN	774	1e-55	90
291	1	700264594H1	SATMON017	g450548	BLASTN	775	1e-55	89
292	1	700074969H1	SATMON007	g450548	BLASTN	777	1e-55	87
293	1	LIB3068-005-Q1-K1-A9	LIB3068	g1778820	BLASTN	389	1e-54	80
294	1	700207147H1	SATMON017	g1778820	BLASTN	537	1e-54	88
295	1	LIB189-004-Q1-E1-F11	LIB189	g960356	BLASTN	757	1e-54	87
296	1	700171222H1	SATMON013	g1778820	BLASTN	757	1e-54	93
297	1	700239903H1	SATMON010	g1778820	BLASTN	760	1e-54	84
298	1	700048208H1	SATMON003	g1778820	BLASTN	762	1e-54	90
299	1	700264085H1	SATMON017	g450548	BLASTN	764	1e-54	87
300	1	700155761H1	SATMON007	g1778820	BLASTN	579	1e-53	85
301	1	700216516H1	SATMON016	g450548	BLASTN	380	1e-52	89
302	1	LIB3068-044-Q1-K1-F12	LIB3068	g450548	BLASTN	492	1e-52	73
303	1	700344420H1	SATMON021	g450548	BLASTN	534	1e-52	79
304	1	700219767H1	SATMON011	g450548	BLASTN	732	1e-52	88
305	1	700153757H1	SATMON007	g1778820	BLASTN	737	1e-52	89
306	1	700165841H1	SATMON013	g450548	BLASTN	738	1e-52	90
307	1	700381966H1	SATMON023	g1778820	BLASTN	740	1e-52	86
308	1	700170427H1	SATMON013	g450548	BLASTN	687	1e-51	89
309	1	700094344H1	SATMON008	g1778820	BLASTN	725	1e-51	93
310	1	700052248H1	SATMON003	g1778820	BLASTN	726	1e-51	84
311	1	700050628H1	SATMON003	g450548	BLASTN	726	1e-51	89
312	1	700223031H1	SATMON011	g1778820	BLASTN	729	1e-51	89
313	1	700475686H1	SATMON025	g450548	BLASTN	660	1e-50	89
314	1	700170325H1	SATMON013	g2305013	BLASTN	709	1e-50	82
315	1	700221107H1	SATMON011	g450548	BLASTN	698	1e-49	89
316	1	700612589H1	SATMON033	g960356	BLASTN	703	1e-49	89
317	1	700153770H1	SATMON007	g1778820	BLASTN	705	1e-49	88
318	1	LIB3078-006-Q1-K1-E7	LIB3078	g1778820	BLASTN	705	1e-49	79

319	1	700239724H1	SATMON010	g1778820	BLASTN	507	1e-48	78
320	1	700356777H1	SATMON024	g1778820	BLASTN	683	1e-48	86
321	1	700241568H1	SATMON010	g960356	BLASTN	691	1e-48	88
322	1	700051294H1	SATMON003	g1778820	BLASTN	462	1e-47	88
323	1	700343661H1	SATMON021	g450548	BLASTN	507	1e-47	81
324	1	700029419H1	SATMON003	g1778820	BLASTN	547	1e-47	87
325	1	700165936H1	SATMON013	g1778820	BLASTN	671	1e-47	83
326	1	700026152H1	SATMON003	g960356	BLASTN	673	1e-47	89
327	1	700075671H1	SATMON007	g450548	BLASTN	429	1e-46	89
328	1	700156674H1	SATMON012	g1778820	BLASTN	659	1e-46	91
329	1	700094981H1	SATMON008	g1778820	BLASTN	662	1e-46	85
330	1	700092879H1	SATMON008	g1778820	BLASTN	668	1e-46	87
331	1	700240685H1	SATMON010	g1778820	BLASTN	484	1e-45	84
332	1	700150286H1	SATMON007	g1778820	BLASTN	647	1e-45	82
333	1	700104990H1	SATMON010	g450548	BLASTN	652	1e-45	89
334	1	700203829H1	SATMON003	g450548	BLASTN	652	1e-45	87
335	1	700153718H1	SATMON007	g167961	BLASTN	656	1e-45	91
336	1	700050841H1	SATMON003	g450548	BLASTN	571	1e-44	86
337	1	700268037H1	SATMON017	g2305013	BLASTN	636	1e-44	86
338	1	700153630H1	SATMON007	g1778820	BLASTN	637	1e-44	82
339	1	700475317H1	SATMON025	g450548	BLASTN	530	1e-43	87
340	1	701163127H1	SATMONN04	g960356	BLASTN	626	1e-43	88
341	1	700203688H1	SATMON003	g960356	BLASTN	627	1e-43	89
342	1	700049893H1	SATMON003	g1778820	BLASTN	506	1e-42	90
343	1	700449155H1	SATMON028	g1778820	BLASTN	443	1e-41	85
344	1	701183780H1	SATMONN06	g450548	BLASTN	558	1e-41	86
345	1	700242162H1	SATMON010	g1778820	BLASTN	600	1e-41	90
346	1	700466994H1	SATMON025	g450548	BLASTN	604	1e-41	91
347	1	700259823H1	SATMON017	g450548	BLASTN	607	1e-41	83
348	1	700346242H1	SATMON021	g450548	BLASTN	608	1e-41	86
349	1	700156395H1	SATMON007	g1778820	BLASTN	609	1e-41	90
350	1	700236835H1	SATMON010	g1778820	BLASTN	335	1e-40	81
351	1	700172385H1	SATMON013	g1778820	BLASTN	397	1e-40	87
352	1	700210466H1	SATMON016	g1778820	BLASTN	586	1e-40	81
353	1	700257303H1	SATMON017	g1778820	BLASTN	589	1e-40	81
354	1	LIB3067-027-Q1-K1-G1	LIB3067	g1778820	BLASTN	623	1e-40	87
355	1	LIB3066-025-Q1-K1-D1	LIB3066	g450548	BLASTN	524	1e-39	85
356	1	700106853H1	SATMON010	g450548	BLASTN	580	1e-39	86
357	1	700160540H1	SATMON012	g960356	BLASTN	581	1e-39	88
358	1	700157780H1	SATMON012	g960356	BLASTN	581	1e-39	88
359	1	700149801H1	SATMON007	g450548	BLASTN	565	1e-38	86
360	1	700353243H1	SATMON024	g450548	BLASTN	570	1e-38	86
361	1	700166171H1	SATMON013	g1778820	BLASTN	254	1e-37	79
362	1	700142509H1	SATMON012	g960356	BLASTN	556	1e-37	88
363	1	700242131H1	SATMON010	g450548	BLASTN	560	1e-37	85
364	1	700455643H1	SATMON029	g450548	BLASTN	539	1e-36	86
365	1	700150248H1	SATMON007	g1778820	BLASTN	547	1e-36	87
366	1	700208549H1	SATMON016	g450548	BLASTN	559	1e-36	88
367	1	700027193H1	SATMON003	g450548	BLASTN	529	1e-35	86
368	1	700221390H1	SATMON011	g450548	BLASTN	530	1e-35	85
369	1	700455647H1	SATMON029	g450548	BLASTN	531	1e-35	85
370	1	700260103H1	SATMON017	g1778820	BLASTN	531	1e-35	80
371	1	700167344H1	SATMON013	g1778820	BLASTN	536	1e-35	88
372	1	700089913H1	SATMON011	g960356	BLASTN	546	1e-35	88

373	1	700570573H1	SATMON030	g450548	BLASTN	300	1e-34	89
374	1	700169889H1	SATMON013	g1778820	BLASTN	519	1e-34	89
375	1	700262857H1	SATMON017	g1778820	BLASTN	521	1e-34	87
376	1	700142644H2	SATMON013	g1778820	BLASTN	522	1e-34	87
377	1	700224417H1	SATMON011	g450548	BLASTN	522	1e-34	91
378	1	700073882H1	SATMON007	g450548	BLASTN	531	1e-34	87
379	1	700085803H1	SATMON011	g450548	BLASTN	338	1e-33	89
380	1	700162323H1	SATMON012	g1778820	BLASTN	513	1e-33	86
381	1	700468306H1	SATMON025	g450548	BLASTN	519	1e-33	89
382	1	700443224H2	SATMON026	g450548	BLASTN	275	1e-32	83
383	1	LIB3066-054-Q1-K1-E3	LIB3066	g450548	BLASTN	495	1e-32	87
384	1	700211827H1	SATMON016	g1778820	BLASTN	489	1e-31	80
385	1	700048741H1	SATMON003	g450548	BLASTN	500	1e-31	88
386	1	700613620H1	SATMON033	g1778820	BLASTN	385	1e-30	88
387	1	700029203H1	SATMON003	g1778820	BLASTN	461	1e-29	84
388	1	700378431H1	SATMON020	g450548	BLASTN	466	1e-29	89
389	1	700455641H1	SATMON029	g450548	BLASTN	472	1e-29	85
390	1	700447867H1	SATMON027	g960356	BLASTN	473	1e-29	88
391	1	700447511H1	SATMON027	g450548	BLASTN	474	1e-29	88
392	1	700025851H1	SATMON003	g450548	BLASTN	479	1e-29	88
393	1	LIB3066-024-Q1-K1-H4	LIB3066	g1778820	BLASTN	481	1e-29	81
394	1	LIB143-025-Q1-E1-C4	LIB143	g450549	BLASTX	64	1e-28	70
395	1	700242282H1	SATMON010	g1778820	BLASTN	280	1e-28	85
396	1	700087244H1	SATMON011	g450548	BLASTN	464	1e-28	88
397	1	700458127H1	SATMON029	g450548	BLASTN	238	1e-27	82
398	1	700025767H1	SATMON003	g1778820	BLASTN	438	1e-26	86
399	1	700235401H1	SATMON010	g450548	BLASTN	442	1e-26	88
400	1	700029457H1	SATMON003	g450548	BLASTN	446	1e-26	89
401	1	700266174H1	SATMON017	g450548	BLASTN	447	1e-26	89
402	1	700349254H1	SATMON023	g1778820	BLASTN	315	1e-25	85
403	1	LIB3068-041-Q1-K1-H6	LIB3068	g450548	BLASTN	438	1e-25	86
404	1	700092889H1	SATMON008	g1778820	BLASTN	397	1e-24	92
405	1	700049044H1	SATMON003	g1778820	BLASTN	425	1e-24	90
406	1	700202710H1	SATMON003	g960357	BLASTX	114	1e-19	97
407	1	700155117H1	SATMON007	g450548	BLASTN	314	1e-19	90
408	1	700449619H1	SATMON028	g1778820	BLASTN	341	1e-19	89
409	1	700150336H1	SATMON007	g1778820	BLASTN	343	1e-19	87
410	1	700166223H1	SATMON013	g960357	BLASTX	181	1e-18	97
411	1	700153796H1	SATMON007	g2305013	BLASTN	300	1e-16	82
412	1	700053266H1	SATMON008	g450548	BLASTN	292	1e-15	83
413	1	700405227H1	SATMON028	g450548	BLASTN	313	1e-15	90
414	1	700397410H1	SATMONN01	g17262	BLASTX	147	1e-14	96
415	1	700211524H1	SATMON016	g1033190	BLASTX	153	1e-14	100
416	1	700281415H2	SATMON019	g2315140	BLASTX	139	1e-13	89
417	1	700429873H1	SATMONN01	g16961	BLASTX	133	1e-12	94
418	1	700239660H1	SATMON010	g450548	BLASTN	245	1e-12	86
419	1	700213596H1	SATMON016	g450548	BLASTN	258	1e-12	95
420	1	700152367H1	SATMON007	g450548	BLASTN	273	1e-12	88
421	1	700452014H1	SATMON028	g17262	BLASTX	76	1e-11	72
422	1	700357106H1	SATMON024	g1724104	BLASTX	132	1e-11	100
423	1	700468611H1	SATMON025	g450549	BLASTX	93	1e-10	93
424	1	700213526H1	SATMON016	g450549	BLASTX	127	1e-10	96
425	1	700267065H1	SATMON017	g450549	BLASTX	130	1e-10	96
426	1	700152044H1	SATMON007	g450549	BLASTX	88	1e-8	93

427	1	700159090H1	SATMON012	g169665	BLASTX	113	1e-8	81
428	1	700266734H1	SATMON017	g450549	BLASTX	119	1e-8	96
429	1	700405367H1	SATMON029	g1778820	BLASTN	231	1e-8	65
1635	-700555532	700555532H1	SOYMON001	g429103	BLASTN	920	1e-67	84
1636	-700649594	700649594H1	SOYMON003	g609559	BLASTX	186	1e-23	85
1637	-700750590	700750590H1	SOYMON014	g609224	BLASTN	363	1e-40	79
1638	-700755802	700755802H1	SOYMON014	g609224	BLASTN	479	1e-43	74
1639	-700869211	700869211H1	SOYMON016	g169665	BLASTX	146	1e-13	92
1640	-700891960	700891960H1	SOYMON024	g726031	BLASTN	589	1e-56	82
1641	-700900377	700900377H1	SOYMON027	g1655577	BLASTN	235	1e-8	78
1642	-700902427	700902427H1	SOYMON027	g1655576	BLASTX	151	1e-13	76
1643	-700941686	700941686H1	SOYMON024	g497899	BLASTN	442	1e-26	89
1644	-700952418	700952418H1	SOYMON022	g726030	BLASTX	146	1e-17	83
1645	-700979651	700979651H2	SOYMON009	g1655579	BLASTN	1006	1e-75	84
1646	-700982809	700982809H1	SOYMON009	g1655579	BLASTN	945	1e-69	82
1647	-700982867	700982867H1	SOYMON009	g1127582	BLASTN	868	1e-63	78
1648	-701056884	701056884H1	SOYMON032	g609556	BLASTN	726	1e-51	84
1649	-701117318	701117318H1	SOYMON037	g609224	BLASTN	451	1e-37	80
1650	-701118224	701118224H1	SOYMON037	g2305013	BLASTN	285	1e-19	72
1651	-701121264	701121264H1	SOYMON037	g166873	BLASTN	238	1e-8	82
1652	-701122908	701122908H1	SOYMON037	g16508	BLASTN	444	1e-26	83
1653	-701128589	701128589H1	SOYMON037	g16508	BLASTN	539	1e-36	90
1654	-GM12798	LIB3049-039-Q1-E1-F2	LIB3049	g16508	BLASTN	578	1e-37	73
1655	-GM14331	LIB3049-055-Q1-E1-F5	LIB3049	g167961	BLASTN	497	1e-30	62
1656	-GM30881	LIB3050-005-Q1-K1-G1	LIB3050	g1655577	BLASTN	543	1e-34	82
1657	-GM30911	LIB3050-005-Q1-K1-B8	LIB3050	g1655577	BLASTN	387	1e-26	71
1658	-GM33921	LIB3051-028-Q1-K1-A9	LIB3051	g16508	BLASTN	338	1e-32	73
1659	12644	701131794H1	SOYMON038	g429107	BLASTN	877	1e-64	84
1660	12644	701142515H1	SOYMON038	g429107	BLASTN	871	1e-63	84
1661	12644	700888494H1	SOYMON024	g429107	BLASTN	662	1e-46	83
1662	16	LIB3030-009-Q1-B1-C1	LIB3030	g429105	BLASTN	1439	1e-119	84
1663	16	LIB3051-106-Q1-K1-B5	LIB3051	g609224	BLASTN	1516	1e-117	86
1664	16	LIB3050-023-Q1-K1-A12	LIB3050	g1724103	BLASTN	1388	1e-106	83
1665	16	LIB3028-003-Q1-B1-G11	LIB3028	g609224	BLASTN	1313	1e-100	87
1666	16	LIB3054-010-Q1-N1-E2	LIB3054	g609224	BLASTN	920	1e-95	87
1667	16	700651294H1	SOYMON003	g609224	BLASTN	672	1e-94	86
1668	16	LIB3027-007-Q1-B1-G3	LIB3027	g16508	BLASTN	891	1e-93	83
1669	16	LIB3053-013-Q1-N1-H9	LIB3053	g609224	BLASTN	996	1e-93	87
1670	16	LIB3051-061-Q1-K1-C7	LIB3051	g16508	BLASTN	1195	1e-93	86
1671	16	LIB3065-005-Q1-N1-A6	LIB3065	g609224	BLASTN	707	1e-91	78
1672	16	LIB3050-008-Q1-E1-E3	LIB3050	g609224	BLASTN	1102	1e-87	87
1673	16	LIB3051-011-Q1-E1-B2	LIB3051	g16508	BLASTN	1153	1e-87	82
1674	16	LIB3030-010-Q1-B1-F8	LIB3030	g609224	BLASTN	988	1e-86	83
1675	16	700652104H1	SOYMON003	g1655577	BLASTN	990	1e-86	83
1676	16	701205279H1	SOYMON035	g609556	BLASTN	1125	1e-85	86
1677	16	700662183H1	SOYMON005	g609224	BLASTN	678	1e-83	84
1678	16	LIB3051-039-Q1-K1-F5	LIB3051	g169664	BLASTN	1104	1e-83	86
1679	16	700557616H1	SOYMON001	g609556	BLASTN	1111	1e-83	86
1680	16	LIB3030-002-Q1-B1-C6	LIB3030	g16508	BLASTN	1113	1e-83	83
1681	16	700865235H1	SOYMON016	g1724103	BLASTN	1090	1e-82	84
1682	16	700563340H1	SOYMON002	g609224	BLASTN	911	1e-80	86
1683	16	LIB3040-044-Q1-E1-D7	LIB3040	g166873	BLASTN	1052	1e-80	82
1684	16	LIB3040-060-Q1-E1-D9	LIB3040	g609224	BLASTN	1071	1e-80	84
1685	16	LIB3049-001-Q1-E1-F12	LIB3049	g16508	BLASTN	1074	1e-80	84

1686	16	LIB3028-006-Q1-B1-F1	LIB3028	g16508	BLASTN	857	1e-79	83
1687	16	LIB3049-033-Q1-E1-G5	LIB3049	g2315139	BLASTN	933	1e-79	81
1688	16	700978240H1	SOYMON009	g609224	BLASTN	984	1e-79	88
1689	16	700980802H1	SOYMON009	g862999	BLASTN	1060	1e-79	85
1690	16	700755908H1	SOYMON014	g1724103	BLASTN	1062	1e-79	88
1691	16	700562226H1	SOYMON002	g1724103	BLASTN	1065	1e-79	84
1692	16	701119101H1	SOYMON037	g609224	BLASTN	1048	1e-78	86
1693	16	700646291H1	SOYMON012	g1655577	BLASTN	1049	1e-78	86
1694	16	LIB3050-022-Q1-K1-B9	LIB3050	g16508	BLASTN	1049	1e-78	83
1695	16	LIB3028-030-Q1-B1-F8	LIB3028	g16508	BLASTN	1053	1e-78	83
1696	16	701143128H1	SOYMON038	g609556	BLASTN	737	1e-77	86
1697	16	LIB3040-041-Q1-E1-D10	LIB3040	g166873	BLASTN	861	1e-77	81
1698	16	700756363H1	SOYMON014	g609556	BLASTN	1031	1e-77	88
1699	16	700729963H1	SOYMON009	g1724103	BLASTN	1036	1e-77	88
1700	16	701063046H1	SOYMON033	g609224	BLASTN	1037	1e-77	86
1701	16	LIB3030-005-Q1-B1-G2	LIB3030	g609224	BLASTN	1038	1e-77	84
1702	16	700564331H1	SOYMON002	g609556	BLASTN	1039	1e-77	84
1703	16	700562958H1	SOYMON002	g1724103	BLASTN	1040	1e-77	88
1704	16	LIB3039-014-Q1-E1-F7	LIB3039	g16508	BLASTN	569	1e-76	84
1705	16	700753632H1	SOYMON014	g497899	BLASTN	668	1e-76	83
1706	16	LIB3049-048-Q1-E1-F2	LIB3049	g16508	BLASTN	722	1e-76	83
1707	16	700648911H1	SOYMON003	g1655577	BLASTN	983	1e-76	83
1708	16	LIB3040-027-Q1-E1-H3	LIB3040	g166873	BLASTN	1003	1e-76	81
1709	16	700664905H1	SOYMON005	g609556	BLASTN	1020	1e-76	86
1710	16	701133404H1	SOYMON038	g1724103	BLASTN	1023	1e-76	85
1711	16	701208780H1	SOYMON035	g1655577	BLASTN	1025	1e-76	85
1712	16	700902279H1	SOYMON027	g169664	BLASTN	1026	1e-76	88
1713	16	LIB3040-048-Q1-E1-G10	LIB3040	g16508	BLASTN	1030	1e-76	83
1714	16	LIB3040-028-Q1-E1-A4	LIB3040	g16508	BLASTN	979	1e-75	84
1715	16	700807586H1	SOYMON016	g609224	BLASTN	1006	1e-75	86
1716	16	700755486H1	SOYMON014	g609224	BLASTN	1006	1e-75	87
1717	16	700724920H1	SOYMON009	g609224	BLASTN	1009	1e-75	87
1718	16	701007494H2	SOYMON019	g1724103	BLASTN	1013	1e-75	85
1719	16	700568310H1	SOYMON002	g497899	BLASTN	834	1e-74	83
1720	16	701208819H1	SOYMON035	g609224	BLASTN	857	1e-74	87
1721	16	700985084H1	SOYMON009	g609224	BLASTN	872	1e-74	85
1722	16	701013074H1	SOYMON019	g726031	BLASTN	894	1e-74	88
1723	16	700650843H1	SOYMON003	g609224	BLASTN	924	1e-74	86
1724	16	700646037H1	SOYMON011	g609556	BLASTN	930	1e-74	84
1725	16	LIB3040-039-Q1-E1-H9	LIB3040	g166873	BLASTN	973	1e-74	80
1726	16	700847231H1	SOYMON021	g726031	BLASTN	999	1e-74	85
1727	16	700792293H1	SOYMON011	g609224	BLASTN	999	1e-74	86
1728	16	701109644H1	SOYMON036	g609556	BLASTN	1004	1e-74	86
1729	16	701122929H1	SOYMON037	g1724103	BLASTN	459	1e-73	87
1730	16	700661491H1	SOYMON005	g16508	BLASTN	540	1e-73	90
1731	16	701120070H1	SOYMON037	g609224	BLASTN	786	1e-73	87
1732	16	700898895H1	SOYMON027	g609556	BLASTN	820	1e-73	88
1733	16	701096959H1	SOYMON028	g429105	BLASTN	839	1e-73	82
1734	16	700848356H1	SOYMON021	g497899	BLASTN	985	1e-73	85
1735	16	700864876H1	SOYMON016	g1724103	BLASTN	985	1e-73	88
1736	16	700868456H1	SOYMON016	g609224	BLASTN	987	1e-73	86
1737	16	701210488H1	SOYMON035	g1655577	BLASTN	988	1e-73	83
1738	16	700747764H1	SOYMON013	g609224	BLASTN	822	1e-72	87
1739	16	701063311H1	SOYMON033	g1655577	BLASTN	971	1e-72	86

1740	16	700868625H1	SOYMON016	g1724103	BLASTN	973	1e-72	85
1741	16	700992344H1	SOYMON011	g726031	BLASTN	973	1e-72	85
1742	16	701012719H1	SOYMON019	g1724103	BLASTN	973	1e-72	85
1743	16	700676769H1	SOYMON007	g609224	BLASTN	973	1e-72	87
1744	16	700946235H1	SOYMON024	g609224	BLASTN	975	1e-72	85
1745	16	700891218H1	SOYMON024	g1724103	BLASTN	975	1e-72	84
1746	16	700724907H1	SOYMON009	g609556	BLASTN	977	1e-72	84
1747	16	700833549H1	SOYMON019	g1655577	BLASTN	979	1e-72	85
1748	16	700942105H1	SOYMON024	g726031	BLASTN	980	1e-72	84
1749	16	700564290H1	SOYMON002	g726031	BLASTN	530	1e-71	84
1750	16	700891233H1	SOYMON024	g1724103	BLASTN	746	1e-71	86
1751	16	LIB3028-025-Q1-B1-B2	LIB3028	g609224	BLASTN	870	1e-71	85
1752	16	701120896H1	SOYMON037	g429105	BLASTN	959	1e-71	82
1753	16	LIB3051-027-Q1-K1-A9	LIB3051	g609224	BLASTN	962	1e-71	83
1754	16	701050696H1	SOYMON032	g609556	BLASTN	965	1e-71	86
1755	16	700653619H1	SOYMON003	g609224	BLASTN	966	1e-71	86
1756	16	701047994H1	SOYMON032	g1724103	BLASTN	967	1e-71	85
1757	16	701123056H1	SOYMON037	g1724103	BLASTN	968	1e-71	85
1758	16	700983479H1	SOYMON009	g726031	BLASTN	653	1e-70	85
1759	16	701063733H1	SOYMON034	g1655577	BLASTN	850	1e-70	85
1760	16	700738366H1	SOYMON012	g726031	BLASTN	946	1e-70	85
1761	16	700866319H1	SOYMON016	g1655575	BLASTN	947	1e-70	85
1762	16	700789013H2	SOYMON011	g1655577	BLASTN	948	1e-70	85
1763	16	700896080H1	SOYMON027	g609224	BLASTN	949	1e-70	87
1764	16	LIB3039-021-Q1-E1-F12	LIB3039	g16508	BLASTN	950	1e-70	83
1765	16	700898284H1	SOYMON027	g429105	BLASTN	950	1e-70	86
1766	16	700901743H1	SOYMON027	g609224	BLASTN	950	1e-70	86
1767	16	700831048H1	SOYMON019	g429105	BLASTN	951	1e-70	85
1768	16	701038107H1	SOYMON029	g726031	BLASTN	952	1e-70	87
1769	16	700559703H1	SOYMON001	g726031	BLASTN	954	1e-70	88
1770	16	700848817H1	SOYMON021	g609224	BLASTN	954	1e-70	84
1771	16	700896032H1	SOYMON027	g609224	BLASTN	956	1e-70	87
1772	16	700944764H1	SOYMON024	g16508	BLASTN	567	1e-69	85
1773	16	701011015H1	SOYMON019	g1724103	BLASTN	647	1e-69	81
1774	16	701125662H1	SOYMON037	g609224	BLASTN	751	1e-69	87
1775	16	701046895H1	SOYMON032	g16508	BLASTN	795	1e-69	83
1776	16	701012632H1	SOYMON019	g1655577	BLASTN	800	1e-69	85
1777	16	701005244H1	SOYMON019	g1724103	BLASTN	804	1e-69	82
1778	16	LIB3050-023-Q1-K1-H10	LIB3050	g609224	BLASTN	899	1e-69	83
1779	16	700892558H1	SOYMON024	g1655575	BLASTN	936	1e-69	85
1780	16	700988779H1	SOYMON011	g16508	BLASTN	937	1e-69	82
1781	16	701203923H2	SOYMON035	g429105	BLASTN	938	1e-69	86
1782	16	701010231H2	SOYMON019	g1724103	BLASTN	938	1e-69	83
1783	16	701041790H1	SOYMON029	g450548	BLASTN	939	1e-69	84
1784	16	700967887H1	SOYMON033	g1724103	BLASTN	940	1e-69	87
1785	16	701123361H1	SOYMON037	g16508	BLASTN	941	1e-69	84
1786	16	701123154H1	SOYMON037	g1724103	BLASTN	942	1e-69	86
1787	16	701045767H1	SOYMON032	g726031	BLASTN	942	1e-69	85
1788	16	700983288H1	SOYMON009	g609224	BLASTN	945	1e-69	83
1789	16	700653053H1	SOYMON003	g609224	BLASTN	945	1e-69	86
1790	16	701044226H1	SOYMON032	g1655577	BLASTN	486	1e-68	83
1791	16	700969927H1	SOYMON005	g497899	BLASTN	750	1e-68	85
1792	16	700547956H1	SOYMON001	g609224	BLASTN	779	1e-68	87
1793	16	LIB3049-048-Q1-E1-E4	LIB3049	g609224	BLASTN	827	1e-68	83

1794	16	701140780H1	SOYMON038	g1655577	BLASTN	922	1e-68	85
1795	16	700849166H1	SOYMON021	g1724103	BLASTN	923	1e-68	84
1796	16	700891945H1	SOYMON024	g609556	BLASTN	925	1e-68	87
1797	16	700658051H1	SOYMON004	g429105	BLASTN	925	1e-68	83
1798	16	700942415H1	SOYMON024	g1655575	BLASTN	928	1e-68	84
1799	16	701041890H1	SOYMON029	g1724103	BLASTN	930	1e-68	82
1800	16	LIB3028-006-Q1-B1-H12	LIB3028	g16508	BLASTN	931	1e-68	81
1801	16	700967039H1	SOYMON029	g429105	BLASTN	931	1e-68	85
1802	16	700836178H1	SOYMON019	g1724103	BLASTN	933	1e-68	85
1803	16	700897558H1	SOYMON027	g16508	BLASTN	737	1e-67	85
1804	16	LIB3050-004-Q1-E1-A2	LIB3050	g609224	BLASTN	796	1e-67	80
1805	16	700730236H1	SOYMON009	g726031	BLASTN	816	1e-67	84
1806	16	700833936H1	SOYMON019	g1724103	BLASTN	915	1e-67	84
1807	16	700897552H1	SOYMON027	g1655577	BLASTN	916	1e-67	84
1808	16	700945440H1	SOYMON024	g609556	BLASTN	917	1e-67	84
1809	16	700961368H1	SOYMON022	g169664	BLASTN	918	1e-67	88
1810	16	700789702H1	SOYMON011	g609224	BLASTN	921	1e-67	86
1811	16	700786541H1	SOYMON011	g429105	BLASTN	513	1e-66	84
1812	16	700987282H1	SOYMON009	g16508	BLASTN	523	1e-66	87
1813	16	700661002H1	SOYMON005	g2305013	BLASTN	836	1e-66	81
1814	16	701148312H1	SOYMON031	g862999	BLASTN	899	1e-66	84
1815	16	700738822H1	SOYMON012	g1655577	BLASTN	899	1e-66	85
1816	16	700940996H1	SOYMON024	g609556	BLASTN	901	1e-66	86
1817	16	700749195H1	SOYMON013	g609556	BLASTN	903	1e-66	81
1818	16	700893941H1	SOYMON024	g429105	BLASTN	903	1e-66	88
1819	16	700892888H1	SOYMON024	g609556	BLASTN	906	1e-66	86
1820	16	700901481H1	SOYMON027	g169664	BLASTN	638	1e-65	86
1821	16	700945269H1	SOYMON024	g169664	BLASTN	688	1e-65	86
1822	16	700746876H1	SOYMON013	g609556	BLASTN	740	1e-65	85
1823	16	700755043H1	SOYMON014	g609556	BLASTN	760	1e-65	87
1824	16	701097166H1	SOYMON028	g1655577	BLASTN	781	1e-65	82
1825	16	701129484H1	SOYMON037	g16508	BLASTN	898	1e-65	82
1826	16	700651014H1	SOYMON003	g167961	BLASTN	464	1e-64	81
1827	16	701134363H1	SOYMON038	g726031	BLASTN	687	1e-64	84
1828	16	701124677H1	SOYMON037	g726031	BLASTN	876	1e-64	85
1829	16	701000359H1	SOYMON018	g1724103	BLASTN	877	1e-64	87
1830	16	701139375H1	SOYMON038	g1724103	BLASTN	883	1e-64	84
1831	16	700832784H1	SOYMON019	g609224	BLASTN	883	1e-64	89
1832	16	700980227H1	SOYMON009	g1724103	BLASTN	884	1e-64	84
1833	16	700943177H1	SOYMON024	g1655577	BLASTN	885	1e-64	85
1834	16	700844446H1	SOYMON021	g2315139	BLASTN	355	1e-63	84
1835	16	700836453H1	SOYMON020	g862999	BLASTN	494	1e-63	85
1836	16	700983012H1	SOYMON009	g16508	BLASTN	756	1e-63	83
1837	16	700795805H1	SOYMON017	g1655577	BLASTN	833	1e-63	85
1838	16	701213663H1	SOYMON035	g1724103	BLASTN	862	1e-63	84
1839	16	700868366H1	SOYMON016	g167961	BLASTN	864	1e-63	83
1840	16	701134377H1	SOYMON038	g2305013	BLASTN	865	1e-63	79
1841	16	LIB3049-007-Q1-E1-C9	LIB3049	g16508	BLASTN	867	1e-63	79
1842	16	700944577H1	SOYMON024	g862999	BLASTN	869	1e-63	85
1843	16	700992289H1	SOYMON011	g16508	BLASTN	869	1e-63	84
1844	16	700891612H1	SOYMON024	g609556	BLASTN	872	1e-63	86
1845	16	700738277H1	SOYMON012	g609224	BLASTN	463	1e-62	85
1846	16	700895571H1	SOYMON027	g609224	BLASTN	562	1e-62	88
1847	16	LIB3049-008-Q1-E1-B3	LIB3049	g16508	BLASTN	853	1e-62	77

1848	16	LIB3027-010-Q1-B1-E12	LIB3027	g609224	BLASTN	854	1e-62	82
1849	16	701129389H1	SOYMON037	g16508	BLASTN	854	1e-62	83
1850	16	701045542H1	SOYMON032	g429105	BLASTN	855	1e-62	86
1851	16	700969901H1	SOYMON005	g726031	BLASTN	856	1e-62	84
1852	16	700984664H1	SOYMON009	g1724103	BLASTN	856	1e-62	81
1853	16	700561352H1	SOYMON002	g609224	BLASTN	858	1e-62	85
1854	16	701138828H1	SOYMON038	g16508	BLASTN	858	1e-62	84
1855	16	700845962H1	SOYMON021	g1655577	BLASTN	861	1e-62	87
1856	16	700896147H1	SOYMON027	g16508	BLASTN	336	1e-61	82
1857	16	701137929H1	SOYMON038	g1655577	BLASTN	502	1e-61	84
1858	16	LIB3040-032-Q1-E1-B8	LIB3040	g16508	BLASTN	521	1e-61	82
1859	16	701009537H1	SOYMON019	g726031	BLASTN	545	1e-61	86
1860	16	LIB3049-031-Q1-E1-E4	LIB3049	g609224	BLASTN	677	1e-61	82
1861	16	LIB3049-031-Q1-E1-C7	LIB3049	g167961	BLASTN	696	1e-61	82
1862	16	700891843H1	SOYMON024	g1655577	BLASTN	737	1e-61	84
1863	16	700561231H1	SOYMON002	g16508	BLASTN	839	1e-61	82
1864	16	700548238H1	SOYMON002	g429105	BLASTN	843	1e-61	85
1865	16	700901018H1	SOYMON027	g2305013	BLASTN	845	1e-61	82
1866	16	701134413H1	SOYMON038	g2305013	BLASTN	848	1e-61	82
1867	16	700653524H1	SOYMON003	g609224	BLASTN	499	1e-60	85
1868	16	LIB3049-017-Q1-E1-G8	LIB3049	g16508	BLASTN	514	1e-60	82
1869	16	701121077H1	SOYMON037	g16508	BLASTN	671	1e-60	84
1870	16	LIB3056-013-Q1-N1-A9	LIB3056	g609224	BLASTN	784	1e-60	86
1871	16	700943524H1	SOYMON024	g16508	BLASTN	831	1e-60	85
1872	16	700952889H1	SOYMON022	g429103	BLASTN	835	1e-60	84
1873	16	700730632H1	SOYMON009	g1655577	BLASTN	431	1e-59	82
1874	16	700649136H1	SOYMON003	g609224	BLASTN	486	1e-59	84
1875	16	700895591H1	SOYMON027	g609556	BLASTN	492	1e-59	87
1876	16	701009829H1	SOYMON019	g429103	BLASTN	572	1e-59	80
1877	16	LIB3039-011-Q1-E1-C11	LIB3039	g16508	BLASTN	619	1e-59	83
1878	16	LIB3040-055-Q1-E1-C4	LIB3040	g16508	BLASTN	622	1e-59	82
1879	16	701123571H1	SOYMON037	g2305013	BLASTN	682	1e-59	81
1880	16	LIB3049-004-Q1-E1-E5	LIB3049	g167961	BLASTN	771	1e-59	83
1881	16	701002239H1	SOYMON018	g609224	BLASTN	782	1e-59	86
1882	16	700971088H1	SOYMON005	g1724103	BLASTN	793	1e-59	81
1883	16	700864624H1	SOYMON016	g167961	BLASTN	817	1e-59	83
1884	16	700863534H1	SOYMON027	g16508	BLASTN	818	1e-59	86
1885	16	700749489H1	SOYMON013	g16508	BLASTN	823	1e-59	86
1886	16	700730689H1	SOYMON009	g16508	BLASTN	825	1e-59	83
1887	16	LIB3049-050-Q1-E1-A11	LIB3049	g16508	BLASTN	831	1e-59	83
1888	16	700850803H1	SOYMON023	g1655577	BLASTN	511	1e-58	83
1889	16	700992317H1	SOYMON011	g1655577	BLASTN	520	1e-58	79
1890	16	701119129H1	SOYMON037	g16508	BLASTN	553	1e-58	83
1891	16	700657623H1	SOYMON004	g1724103	BLASTN	600	1e-58	83
1892	16	LIB3049-015-Q1-E1-F8	LIB3049	g16508	BLASTN	674	1e-58	79
1893	16	701007027H1	SOYMON019	g16508	BLASTN	804	1e-58	81
1894	16	701120820H1	SOYMON037	g609224	BLASTN	807	1e-58	85
1895	16	700654317H1	SOYMON004	g1655577	BLASTN	811	1e-58	84
1896	16	700889071H1	SOYMON024	g497899	BLASTN	812	1e-58	83
1897	16	701010676H1	SOYMON019	g609224	BLASTN	813	1e-58	85
1898	16	701099590H1	SOYMON028	g1655577	BLASTN	486	1e-57	82
1899	16	700649684H1	SOYMON003	g16508	BLASTN	540	1e-57	82
1900	16	700994107H1	SOYMON011	g16508	BLASTN	567	1e-57	80
1901	16	700898629H1	SOYMON027	g16508	BLASTN	687	1e-57	82

1902	16	700902333H1	SOYMON027	g609224	BLASTN	700	1e-57	82
1903	16	LIB3039-017-Q1-E1-D9	LIB3039	g16508	BLASTN	703	1e-57	81
1904	16	LIB3040-026-Q1-E1-A4	LIB3040	g16508	BLASTN	709	1e-57	83
1905	16	701130101H1	SOYMON037	g1724103	BLASTN	791	1e-57	88
1906	16	700902482H1	SOYMON027	g169664	BLASTN	797	1e-57	81
1907	16	700730789H1	SOYMON009	g16508	BLASTN	798	1e-57	82
1908	16	700868670H1	SOYMON016	g16508	BLASTN	800	1e-57	83
1909	16	LIB3029-011-Q1-B1-D1	LIB3029	g609224	BLASTN	800	1e-57	82
1910	16	701119278H1	SOYMON037	g609224	BLASTN	801	1e-57	83
1911	16	700747731H1	SOYMON013	g16508	BLASTN	801	1e-57	86
1912	16	LIB3049-017-Q1-E1-A11	LIB3049	g16508	BLASTN	811	1e-57	80
1913	16	700890428H1	SOYMON024	g609224	BLASTN	482	1e-56	83
1914	16	700562326H1	SOYMON002	g609224	BLASTN	778	1e-56	82
1915	16	700567740H1	SOYMON002	g609224	BLASTN	781	1e-56	85
1916	16	701061514H1	SOYMON033	g1724103	BLASTN	784	1e-56	86
1917	16	701135670H1	SOYMON038	g429105	BLASTN	785	1e-56	81
1918	16	701139657H1	SOYMON038	g609224	BLASTN	786	1e-56	85
1919	16	701123371H1	SOYMON037	g16508	BLASTN	788	1e-56	86
1920	16	700838744H1	SOYMON020	g16508	BLASTN	789	1e-56	82
1921	16	701070458H1	SOYMON034	g1655577	BLASTN	457	1e-55	78
1922	16	701003437H1	SOYMON019	g429105	BLASTN	474	1e-55	83
1923	16	700835976H1	SOYMON019	g16508	BLASTN	562	1e-55	84
1924	16	700894535H1	SOYMON024	g16508	BLASTN	570	1e-55	86
1925	16	700752755H1	SOYMON014	g1655577	BLASTN	690	1e-55	83
1926	16	LIB3049-032-Q1-E1-C9	LIB3049	g16508	BLASTN	700	1e-55	83
1927	16	LIB3040-030-Q1-E1-B5	LIB3040	g16508	BLASTN	709	1e-55	84
1928	16	701136935H1	SOYMON038	g609224	BLASTN	766	1e-55	85
1929	16	700682088H1	SOYMON008	g169664	BLASTN	767	1e-55	90
1930	16	700682188H1	SOYMON008	g169664	BLASTN	767	1e-55	90
1931	16	701068649H1	SOYMON034	g16508	BLASTN	770	1e-55	83
1932	16	700726623H1	SOYMON009	g16508	BLASTN	771	1e-55	85
1933	16	700751129H1	SOYMON014	g16508	BLASTN	772	1e-55	86
1934	16	700945234H1	SOYMON024	g16508	BLASTN	773	1e-55	83
1935	16	701133507H2	SOYMON038	g609224	BLASTN	776	1e-55	85
1936	16	700902459H1	SOYMON027	g726031	BLASTN	777	1e-55	84
1937	16	700986624H1	SOYMON009	g16508	BLASTN	612	1e-54	83
1938	16	701097020H1	SOYMON028	g609224	BLASTN	615	1e-54	84
1939	16	701131409H1	SOYMON038	g16508	BLASTN	755	1e-54	91
1940	16	700750123H1	SOYMON013	g726031	BLASTN	756	1e-54	86
1941	16	701040251H1	SOYMON029	g16508	BLASTN	756	1e-54	85
1942	16	700732290H1	SOYMON010	g169664	BLASTN	758	1e-54	90
1943	16	701117458H1	SOYMON037	g609224	BLASTN	759	1e-54	82
1944	16	701118709H1	SOYMON037	g16508	BLASTN	760	1e-54	91
1945	16	701012834H1	SOYMON019	g16508	BLASTN	760	1e-54	91
1946	16	701133316H1	SOYMON038	g16508	BLASTN	760	1e-54	91
1947	16	701129646H1	SOYMON037	g16508	BLASTN	760	1e-54	91
1948	16	700732265H1	SOYMON010	g169664	BLASTN	760	1e-54	90
1949	16	701040796H1	SOYMON029	g1724103	BLASTN	760	1e-54	89
1950	16	700846350H1	SOYMON021	g16508	BLASTN	761	1e-54	86
1951	16	701137005H1	SOYMON038	g16508	BLASTN	761	1e-54	86
1952	16	701046506H1	SOYMON032	g16508	BLASTN	761	1e-54	91
1953	16	700894462H1	SOYMON024	g16508	BLASTN	761	1e-54	86
1954	16	700973847H1	SOYMON005	g16508	BLASTN	766	1e-54	87
1955	16	701128215H1	SOYMON037	g609556	BLASTN	466	1e-53	82

1956	16	700842468H1	SOYMON020	g429105	BLASTN	548	1e-53	86
1957	16	701051552H1	SOYMON032	g609224	BLASTN	744	1e-53	85
1958	16	700944049H1	SOYMON024	g609224	BLASTN	744	1e-53	85
1959	16	701120261H1	SOYMON037	g609224	BLASTN	746	1e-53	84
1960	16	700897524H1	SOYMON027	g609224	BLASTN	751	1e-53	84
1961	16	LIB3049-042-Q1-E1-F7	LIB3049	g450548	BLASTN	761	1e-53	75
1962	16	700896909H1	SOYMON027	g429105	BLASTN	536	1e-52	82
1963	16	700974820H1	SOYMON005	g1724103	BLASTN	582	1e-52	86
1964	16	701143224H1	SOYMON038	g609224	BLASTN	731	1e-52	84
1965	16	701010322H1	SOYMON019	g609224	BLASTN	731	1e-52	85
1966	16	701130510H1	SOYMON038	g609224	BLASTN	732	1e-52	83
1967	16	700724904H1	SOYMON009	g609224	BLASTN	733	1e-52	85
1968	16	701119845H1	SOYMON037	g609224	BLASTN	734	1e-52	85
1969	16	701107871H1	SOYMON036	g16508	BLASTN	737	1e-52	86
1970	16	700983380H1	SOYMON009	g609224	BLASTN	738	1e-52	84
1971	16	700896469H1	SOYMON027	g16508	BLASTN	741	1e-52	86
1972	16	700792178H1	SOYMON011	g16508	BLASTN	742	1e-52	85
1973	16	701099906H1	SOYMON028	g16508	BLASTN	431	1e-51	89
1974	16	701134724H2	SOYMON038	g16508	BLASTN	553	1e-51	85
1975	16	700749712H1	SOYMON013	g16508	BLASTN	689	1e-51	87
1976	16	700982567H1	SOYMON009	g609224	BLASTN	723	1e-51	85
1977	16	701013758H1	SOYMON019	g609224	BLASTN	723	1e-51	85
1978	16	700868508H1	SOYMON016	g609224	BLASTN	723	1e-51	85
1979	16	700749316H1	SOYMON013	g609224	BLASTN	723	1e-51	85
1980	16	701045367H1	SOYMON032	g16508	BLASTN	724	1e-51	91
1981	16	701205494H1	SOYMON035	g609224	BLASTN	724	1e-51	84
1982	16	700556784H1	SOYMON001	g16508	BLASTN	724	1e-51	91
1983	16	701099879H1	SOYMON028	g16508	BLASTN	726	1e-51	80
1984	16	701131671H1	SOYMON038	g609224	BLASTN	728	1e-51	85
1985	16	701011505H1	SOYMON019	g609224	BLASTN	728	1e-51	85
1986	16	701009973H2	SOYMON019	g609224	BLASTN	728	1e-51	85
1987	16	700793520H1	SOYMON017	g609224	BLASTN	728	1e-51	85
1988	16	700753622H1	SOYMON014	g609224	BLASTN	728	1e-51	85
1989	16	700984052H1	SOYMON009	g609224	BLASTN	728	1e-51	83
1990	16	701108521H1	SOYMON036	g609224	BLASTN	728	1e-51	85
1991	16	700957203H1	SOYMON022	g1724103	BLASTN	464	1e-50	86
1992	16	700554120H1	SOYMON001	g609224	BLASTN	528	1e-50	84
1993	16	700555954H1	SOYMON001	g167961	BLASTN	528	1e-50	86
1994	16	701124141H1	SOYMON037	g609224	BLASTN	537	1e-50	85
1995	16	701003232H1	SOYMON019	g16508	BLASTN	544	1e-50	88
1996	16	700653112H1	SOYMON003	g16508	BLASTN	559	1e-50	91
1997	16	701103353H1	SOYMON028	g1655577	BLASTN	581	1e-50	84
1998	16	700562790H1	SOYMON002	g497899	BLASTN	708	1e-50	86
1999	16	701097303H1	SOYMON028	g609224	BLASTN	711	1e-50	84
2000	16	700561195H1	SOYMON002	g609224	BLASTN	712	1e-50	84
2001	16	701121162H1	SOYMON037	g16508	BLASTN	712	1e-50	88
2002	16	700981317H1	SOYMON009	g16508	BLASTN	714	1e-50	89
2003	16	700981595H1	SOYMON009	g16508	BLASTN	716	1e-50	86
2004	16	700994305H1	SOYMON011	g609224	BLASTN	528	1e-49	85
2005	16	701101565H1	SOYMON028	g429105	BLASTN	552	1e-49	82
2006	16	701103456H1	SOYMON028	g429105	BLASTN	617	1e-49	81
2007	16	700993331H1	SOYMON011	g16508	BLASTN	622	1e-49	86
2008	16	LIB3052-011-Q1-N1-B12	LIB3052	g16508	BLASTN	695	1e-49	85
2009	16	701140613H1	SOYMON038	g609224	BLASTN	697	1e-49	85

2010	16	700745426H1	SOYMON013	g16508	BLASTN	697	1e-49	84
2011	16	701046530H1	SOYMON032	g609224	BLASTN	697	1e-49	84
2012	16	701003787H1	SOYMON019	g609224	BLASTN	697	1e-49	85
2013	16	700840830H1	SOYMON020	g1724103	BLASTN	699	1e-49	79
2014	16	700901588H1	SOYMON027	g609224	BLASTN	702	1e-49	85
2015	16	701037824H1	SOYMON029	g497899	BLASTN	702	1e-49	87
2016	16	701131888H1	SOYMON038	g609224	BLASTN	702	1e-49	85
2017	16	701009947H2	SOYMON019	g609224	BLASTN	702	1e-49	85
2018	16	700729216H1	SOYMON009	g497899	BLASTN	702	1e-49	87
2019	16	700831705H1	SOYMON019	g609224	BLASTN	702	1e-49	85
2020	16	700900329H1	SOYMON027	g429105	BLASTN	705	1e-49	88
2021	16	700563946H1	SOYMON002	g16508	BLASTN	705	1e-49	83
2022	16	700808370H1	SOYMON024	g609556	BLASTN	459	1e-48	83
2023	16	LIB3040-001-Q1-E1-H4	LIB3040	g1724103	BLASTN	475	1e-48	76
2024	16	700989482H1	SOYMON011	g16508	BLASTN	498	1e-48	86
2025	16	701001311H1	SOYMON018	g16508	BLASTN	655	1e-48	85
2026	16	700734733H1	SOYMON010	g497899	BLASTN	682	1e-48	87
2027	16	701013070H1	SOYMON019	g497899	BLASTN	682	1e-48	87
2028	16	701120989H1	SOYMON037	g16508	BLASTN	683	1e-48	91
2029	16	LIB3049-025-Q1-E1-B4	LIB3049	g167961	BLASTN	687	1e-48	82
2030	16	700808455H1	SOYMON024	g16508	BLASTN	688	1e-48	87
2031	16	701101319H1	SOYMON028	g16508	BLASTN	688	1e-48	91
2032	16	701037087H1	SOYMON029	g609224	BLASTN	689	1e-48	85
2033	16	701205225H1	SOYMON035	g16508	BLASTN	689	1e-48	84
2034	16	701136807H1	SOYMON038	g609224	BLASTN	690	1e-48	85
2035	16	701118459H1	SOYMON037	g609224	BLASTN	690	1e-48	85
2036	16	700942825H1	SOYMON024	g16508	BLASTN	691	1e-48	85
2037	16	701139796H1	SOYMON038	g497899	BLASTN	692	1e-48	87
2038	16	700557040H1	SOYMON001	g609224	BLASTN	692	1e-48	85
2039	16	701015715H1	SOYMON038	g497899	BLASTN	692	1e-48	87
2040	16	700726424H1	SOYMON009	g497899	BLASTN	693	1e-48	87
2041	16	700790811H1	SOYMON011	g497899	BLASTN	693	1e-48	87
2042	16	LIB3040-038-Q1-E1-E5	LIB3040	g16508	BLASTN	709	1e-48	85
2043	16	700896626H1	SOYMON027	g429107	BLASTN	430	1e-47	84
2044	16	700562158H1	SOYMON002	g609224	BLASTN	518	1e-47	84
2045	16	700747095H1	SOYMON013	g16508	BLASTN	538	1e-47	87
2046	16	700726532H1	SOYMON009	g609224	BLASTN	622	1e-47	85
2047	16	701137686H1	SOYMON038	g497899	BLASTN	671	1e-47	85
2048	16	701009148H1	SOYMON019	g16508	BLASTN	673	1e-47	91
2049	16	701048279H1	SOYMON032	g16508	BLASTN	674	1e-47	86
2050	16	701120185H1	SOYMON037	g16508	BLASTN	674	1e-47	91
2051	16	701040355H1	SOYMON029	g497899	BLASTN	675	1e-47	86
2052	16	701130203H1	SOYMON037	g609224	BLASTN	678	1e-47	83
2053	16	701015794H1	SOYMON038	g16508	BLASTN	681	1e-47	91
2054	16	700732181H1	SOYMON010	g16508	BLASTN	681	1e-47	86
2055	16	LIB3050-018-Q1-E1-D10	LIB3050	g2305013	BLASTN	696	1e-47	81
2056	16	700899157H1	SOYMON027	g609224	BLASTN	486	1e-46	84
2057	16	701119905H1	SOYMON037	g497899	BLASTN	507	1e-46	86
2058	16	701062873H1	SOYMON033	g497899	BLASTN	509	1e-46	88
2059	16	701210886H1	SOYMON035	g16508	BLASTN	540	1e-46	84
2060	16	700893278H1	SOYMON024	g16508	BLASTN	542	1e-46	77
2061	16	701036970H1	SOYMON029	g16508	BLASTN	579	1e-46	88
2062	16	700901932H1	SOYMON027	g609224	BLASTN	640	1e-46	85
2063	16	701044214H1	SOYMON032	g429105	BLASTN	658	1e-46	85

2064	16	701056917H1	SOYMON033	g16508	BLASTN	661	1e-46	85
2065	16	700981560H1	SOYMON009	g497899	BLASTN	661	1e-46	87
2066	16	701213107H1	SOYMON035	g609224	BLASTN	661	1e-46	83
2067	16	700834235H1	SOYMON019	g497899	BLASTN	661	1e-46	87
2068	16	700749413H1	SOYMON013	g16508	BLASTN	662	1e-46	85
2069	16	700889148H1	SOYMON024	g16508	BLASTN	662	1e-46	85
2070	16	700745009H1	SOYMON013	g609224	BLASTN	662	1e-46	85
2071	16	701206618H1	SOYMON035	g16508	BLASTN	663	1e-46	91
2072	16	701131927H1	SOYMON038	g497899	BLASTN	666	1e-46	87
2073	16	701119069H1	SOYMON037	g497899	BLASTN	666	1e-46	87
2074	16	701106908H1	SOYMON036	g497899	BLASTN	666	1e-46	87
2075	16	701103063H1	SOYMON028	g497899	BLASTN	666	1e-46	87
2076	16	700686659H1	SOYMON008	g166873	BLASTN	667	1e-46	86
2077	16	701040620H1	SOYMON029	g16508	BLASTN	669	1e-46	86
2078	16	701012134H1	SOYMON019	g497899	BLASTN	669	1e-46	85
2079	16	700957508H1	SOYMON022	g1724103	BLASTN	669	1e-46	82
2080	16	701008548H1	SOYMON019	g497899	BLASTN	500	1e-45	87
2081	16	700978303H1	SOYMON009	g497899	BLASTN	506	1e-45	87
2082	16	701119990H1	SOYMON037	g497899	BLASTN	512	1e-45	87
2083	16	700646222H1	SOYMON012	g16508	BLASTN	526	1e-45	85
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2085	16	700894009H1	SOYMON024	g497899	BLASTN	646	1e-45	87
2086	16	700908702H1	SOYMON022	g609224	BLASTN	646	1e-45	85
2087	16	700838642H1	SOYMON020	g609224	BLASTN	646	1e-45	85
2088	16	700832089H1	SOYMON019	g609224	BLASTN	646	1e-45	85
2089	16	700978279H1	SOYMON009	g609224	BLASTN	647	1e-45	80
2090	16	700555058H1	SOYMON001	g16508	BLASTN	649	1e-45	84
2091	16	700906478H1	SOYMON022	g497899	BLASTN	651	1e-45	87
2092	16	700897237H1	SOYMON027	g16508	BLASTN	653	1e-45	86
2093	16	701125786H1	SOYMON037	g16508	BLASTN	654	1e-45	84
2094	16	700562836H1	SOYMON002	g16508	BLASTN	655	1e-45	84
2095	16	700665969H1	SOYMON005	g497899	BLASTN	656	1e-45	87
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2100	16	701048175H1	SOYMON032	g16508	BLASTN	658	1e-45	91
2101	16	700901037H1	SOYMON027	g16508	BLASTN	658	1e-45	86
2102	16	700656917H1	SOYMON004	g1724103	BLASTN	407	1e-44	81
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2104	16	701002808H1	SOYMON019	g497899	BLASTN	503	1e-44	87
2105	16	701107115H1	SOYMON036	g497899	BLASTN	507	1e-44	87
2106	16	701122669H1	SOYMON037	g16508	BLASTN	636	1e-44	82
2107	16	700894409H1	SOYMON024	g497899	BLASTN	636	1e-44	87
2108	16	700848374H1	SOYMON021	g497899	BLASTN	636	1e-44	87
2109	16	700902420H1	SOYMON027	g497899	BLASTN	636	1e-44	87
2110	16	700747450H1	SOYMON013	g16508	BLASTN	638	1e-44	91
2111	16	701125869H1	SOYMON037	g16508	BLASTN	638	1e-44	91
2112	16	700731302H1	SOYMON010	g16508	BLASTN	639	1e-44	85
2113	16	701099068H1	SOYMON028	g2305013	BLASTN	640	1e-44	81
2114	16	700975556H1	SOYMON009	g450548	BLASTN	641	1e-44	84
2115	16	700889660H1	SOYMON024	g16508	BLASTN	642	1e-44	86
2116	16	700951744H1	SOYMON022	g16508	BLASTN	642	1e-44	86
2117	16	700750251H1	SOYMON013	g609224	BLASTN	643	1e-44	86

2118	16	700834611H1	SOYMON019	g16508	BLASTN	643	1e-44	91
2119	16	700565356H1	SOYMON002	g16508	BLASTN	643	1e-44	80
2120	16	700673803H1	SOYMON007	g16508	BLASTN	644	1e-44	84
2121	16	700958721H1	SOYMON022	g16508	BLASTN	645	1e-44	86
2122	16	700985717H1	SOYMON009	g16508	BLASTN	646	1e-44	82
2123	16	LIB3049-032-Q1-E1-A5	LIB3049	g16508	BLASTN	662	1e-44	76
2124	16	700733454H1	SOYMON010	g609224	BLASTN	432	1e-43	85
2125	16	700987656H1	SOYMON009	g497899	BLASTN	476	1e-43	87
2126	16	700755957H1	SOYMON014	g497899	BLASTN	483	1e-43	86
2127	16	700749331H1	SOYMON013	g2305013	BLASTN	521	1e-43	84
2128	16	700830984H1	SOYMON019	g497899	BLASTN	622	1e-43	86
2129	16	700906176H1	SOYMON022	g16508	BLASTN	623	1e-43	86
2130	16	701100070H2	SOYMON028	g16508	BLASTN	623	1e-43	91
2131	16	700954053H1	SOYMON022	g497899	BLASTN	624	1e-43	88
2132	16	700833949H1	SOYMON019	g497899	BLASTN	624	1e-43	88
2133	16	700976710H1	SOYMON009	g2305013	BLASTN	625	1e-43	79
2134	16	701117925H2	SOYMON037	g16508	BLASTN	625	1e-43	84
2135	16	700748825H1	SOYMON013	g16508	BLASTN	626	1e-43	84
2136	16	700899651H1	SOYMON027	g16508	BLASTN	626	1e-43	84
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2138	16	701009051H1	SOYMON019	g16508	BLASTN	628	1e-43	91
2139	16	700754423H1	SOYMON014	g16508	BLASTN	628	1e-43	91
2140	16	700760701H1	SOYMON015	g16508	BLASTN	628	1e-43	83
2141	16	701098124H1	SOYMON028	g2305013	BLASTN	628	1e-43	78
2142	16	700984979H1	SOYMON009	g16508	BLASTN	628	1e-43	91
2143	16	700565262H1	SOYMON002	g497899	BLASTN	629	1e-43	86
2144	16	700954979H1	SOYMON022	g497899	BLASTN	630	1e-43	87
2145	16	700834601H1	SOYMON019	g497899	BLASTN	630	1e-43	87
2146	16	700953156H1	SOYMON022	g609224	BLASTN	630	1e-43	85
2147	16	700865217H1	SOYMON016	g16508	BLASTN	631	1e-43	77
2148	16	700852947H1	SOYMON023	g16508	BLASTN	631	1e-43	84
2149	16	700943547H1	SOYMON024	g497899	BLASTN	631	1e-43	86
2150	16	701139277H1	SOYMON038	g609224	BLASTN	632	1e-43	82
2151	16	700900155H1	SOYMON027	g429105	BLASTN	264	1e-42	81
2152	16	700970581H1	SOYMON005	g497899	BLASTN	296	1e-42	86
2153	16	700654232H1	SOYMON003	g497899	BLASTN	348	1e-42	88
2154	16	700743608H1	SOYMON012	g497899	BLASTN	354	1e-42	86
2155	16	700986620H1	SOYMON009	g609224	BLASTN	360	1e-42	84
2156	16	700562590H1	SOYMON002	g16508	BLASTN	476	1e-42	83
2157	16	701119690H1	SOYMON037	g167961	BLASTN	506	1e-42	84
2158	16	700951730H1	SOYMON022	g497899	BLASTN	507	1e-42	87
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2163	16	700547973H1	SOYMON001	g497899	BLASTN	611	1e-42	84
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2166	16	700946136H1	SOYMON024	g16508	BLASTN	615	1e-42	85
2167	16	700665932H1	SOYMON005	g16508	BLASTN	616	1e-42	84
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2169	16	700962515H1	SOYMON022	g16508	BLASTN	618	1e-42	91
2170	16	700895685H1	SOYMON027	g429107	BLASTN	618	1e-42	77
2171	16	700752015H1	SOYMON014	g16508	BLASTN	620	1e-42	84

2172	16	701015704H1	SOYMON038	g16508	BLASTN	620	1e-42	85
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2178	16	700555653H1	SOYMON001	g166873	BLASTN	512	1e-41	85
2179	16	700958314H1	SOYMON022	g497899	BLASTN	599	1e-41	87
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2181	16	700901925H1	SOYMON027	g16508	BLASTN	599	1e-41	84
2182	16	700741987H1	SOYMON012	g497899	BLASTN	599	1e-41	87
2183	16	700662837H1	SOYMON005	g497899	BLASTN	599	1e-41	87
2184	16	701060927H1	SOYMON033	g497899	BLASTN	599	1e-41	87
2185	16	700562973H1	SOYMON002	g16508	BLASTN	601	1e-41	87
2186	16	700754586H1	SOYMON014	g16508	BLASTN	603	1e-41	84
2187	16	701132674H1	SOYMON038	g16508	BLASTN	604	1e-41	85
2188	16	701010012H2	SOYMON019	g609224	BLASTN	604	1e-41	84
2189	16	701123643H1	SOYMON037	g16508	BLASTN	604	1e-41	85
2190	16	700894196H1	SOYMON024	g16508	BLASTN	604	1e-41	85
2191	16	700899248H1	SOYMON027	g16508	BLASTN	604	1e-41	85
2192	16	701207680H1	SOYMON035	g16508	BLASTN	605	1e-41	84
2193	16	700899210H1	SOYMON027	g16508	BLASTN	605	1e-41	84
2194	16	700898762H1	SOYMON027	g16508	BLASTN	605	1e-41	87
2195	16	700983636H1	SOYMON009	g497899	BLASTN	606	1e-41	87
2196	16	700845511H1	SOYMON021	g16508	BLASTN	606	1e-41	89
2197	16	700563416H1	SOYMON002	g16508	BLASTN	606	1e-41	83
2198	16	701004041H1	SOYMON019	g16508	BLASTN	607	1e-41	84
2199	16	701097184H1	SOYMON028	g16508	BLASTN	607	1e-41	87
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2203	16	700831972H1	SOYMON019	g16508	BLASTN	609	1e-41	85
2204	16	701010090H2	SOYMON019	g16508	BLASTN	610	1e-41	87
2205	16	700890067H1	SOYMON024	g16508	BLASTN	610	1e-41	87
2206	16	700833227H1	SOYMON019	g16508	BLASTN	610	1e-41	85
2207	16	700563171H1	SOYMON002	g16508	BLASTN	370	1e-40	90
2208	16	701121457H1	SOYMON037	g609224	BLASTN	372	1e-40	85
2209	16	700566851H1	SOYMON002	g16508	BLASTN	402	1e-40	82
2210	16	700753221H1	SOYMON014	g609224	BLASTN	422	1e-40	84
2211	16	700749039H1	SOYMON013	g16508	BLASTN	475	1e-40	84
2212	16	700558940H1	SOYMON001	g16508	BLASTN	493	1e-40	80
2213	16	700757695H1	SOYMON015	g16508	BLASTN	526	1e-40	83
2214	16	701102262H1	SOYMON028	g2305013	BLASTN	587	1e-40	81
2215	16	700747484H1	SOYMON013	g16508	BLASTN	587	1e-40	85
2216	16	700891875H1	SOYMON024	g16508	BLASTN	587	1e-40	85
2217	16	700894439H1	SOYMON024	g497899	BLASTN	588	1e-40	88
2218	16	701062577H1	SOYMON033	g16508	BLASTN	588	1e-40	87
2219	16	700964366H1	SOYMON022	g16508	BLASTN	589	1e-40	85
2220	16	701210016H1	SOYMON035	g16508	BLASTN	589	1e-40	83
2221	16	700901573H1	SOYMON027	g16508	BLASTN	589	1e-40	85
2222	16	700750533H1	SOYMON014	g16508	BLASTN	589	1e-40	85
2223	16	700888820H1	SOYMON024	g497899	BLASTN	589	1e-40	87
2224	16	700891965H1	SOYMON024	g497899	BLASTN	589	1e-40	87
2225	16	700865001H1	SOYMON016	g16508	BLASTN	589	1e-40	85

2226	16	700654923H1	SOYMON004	g16508	BLASTN	590	1e-40	84
2227	16	700746169H1	SOYMON013	g16508	BLASTN	590	1e-40	84
2228	16	700745792H1	SOYMON013	g16508	BLASTN	591	1e-40	85
2229	16	700966953H1	SOYMON029	g16508	BLASTN	591	1e-40	85
2230	16	701141657H1	SOYMON038	g16508	BLASTN	592	1e-40	86
2231	16	700943078H1	SOYMON024	g16508	BLASTN	598	1e-40	85
2232	16	701137073H1	SOYMON038	g16508	BLASTN	598	1e-40	90
2233	16	700966730H1	SOYMON028	g16508	BLASTN	598	1e-40	84
2234	16	701137731H1	SOYMON038	g16508	BLASTN	371	1e-39	84
2235	16	701044050H1	SOYMON032	g497899	BLASTN	428	1e-39	89
2236	16	700993309H1	SOYMON011	g497899	BLASTN	434	1e-39	88
2237	16	700874483H1	SOYMON018	g497899	BLASTN	434	1e-39	87
2238	16	701014866H1	SOYMON019	g16508	BLASTN	454	1e-39	89
2239	16	700894603H1	SOYMON024	g16508	BLASTN	454	1e-39	85
2240	16	700958677H1	SOYMON022	g497899	BLASTN	455	1e-39	87
2241	16	700754001H1	SOYMON014	g609224	BLASTN	518	1e-39	85
2242	16	700946430H1	SOYMON024	g497899	BLASTN	578	1e-39	87
2243	16	700729892H1	SOYMON009	g497899	BLASTN	578	1e-39	87
2244	16	700753230H1	SOYMON014	g497899	BLASTN	578	1e-39	87
2245	16	700562584H1	SOYMON002	g16508	BLASTN	578	1e-39	82
2246	16	700842880H1	SOYMON020	g16508	BLASTN	580	1e-39	82
2247	16	700852561H1	SOYMON023	g16508	BLASTN	582	1e-39	85
2248	16	700941290H1	SOYMON024	g497899	BLASTN	583	1e-39	87
2249	16	700750361H1	SOYMON013	g497899	BLASTN	584	1e-39	80
2250	16	700841915H1	SOYMON020	g16508	BLASTN	585	1e-39	85
2251	16	700944462H1	SOYMON024	g16508	BLASTN	585	1e-39	85
2252	16	700548174H1	SOYMON002	g16508	BLASTN	342	1e-38	82
2253	16	700942324H1	SOYMON024	g497899	BLASTN	403	1e-38	87
2254	16	700788450H1	SOYMON011	g497899	BLASTN	416	1e-38	86
2255	16	701127954H1	SOYMON037	g16508	BLASTN	449	1e-38	76
2256	16	700742223H1	SOYMON012	g1724103	BLASTN	451	1e-38	77
2257	16	701102722H1	SOYMON028	g16508	BLASTN	466	1e-38	92
2258	16	701046957H1	SOYMON032	g16508	BLASTN	517	1e-38	85
2259	16	700836169H1	SOYMON019	g497899	BLASTN	517	1e-38	87
2260	16	700697967H1	SOYMON015	g16508	BLASTN	521	1e-38	84
2261	16	700678867H1	SOYMON007	g609224	BLASTN	566	1e-38	82
2262	16	700838753H1	SOYMON020	g16508	BLASTN	567	1e-38	85
2263	16	700726468H1	SOYMON009	g16508	BLASTN	570	1e-38	85
2264	16	701014631H1	SOYMON019	g497899	BLASTN	571	1e-38	87
2265	16	700728719H1	SOYMON009	g497899	BLASTN	573	1e-38	87
2266	16	700962582H1	SOYMON022	g497899	BLASTN	573	1e-38	87
2267	16	701123183H1	SOYMON037	g16508	BLASTN	573	1e-38	84
2268	16	700752347H1	SOYMON014	g16508	BLASTN	574	1e-38	87
2269	16	701103315H1	SOYMON028	g16508	BLASTN	394	1e-37	86
2270	16	700750955H1	SOYMON014	g497899	BLASTN	414	1e-37	88
2271	16	700867493H1	SOYMON016	g609224	BLASTN	433	1e-37	85
2272	16	701054760H1	SOYMON032	g16508	BLASTN	475	1e-37	85
2273	16	700900903H1	SOYMON027	g16508	BLASTN	501	1e-37	84
2274	16	701102294H1	SOYMON028	g16508	BLASTN	502	1e-37	87
2275	16	701124034H1	SOYMON037	g16508	BLASTN	512	1e-37	82
2276	16	700981476H1	SOYMON009	g16508	BLASTN	526	1e-37	83
2277	16	700903712H1	SOYMON022	g609224	BLASTN	552	1e-37	85
2278	16	700648751H1	SOYMON003	g1655577	BLASTN	554	1e-37	86
2279	16	700829930H1	SOYMON019	g609224	BLASTN	556	1e-37	84

2280	16	700758078H1	SOYMON015	g16508	BLASTN	556	1e-37	84
2281	16	700754248H1	SOYMON014	g16508	BLASTN	557	1e-37	81
2282	16	700834650H1	SOYMON019	g609224	BLASTN	557	1e-37	85
2283	16	700746279H1	SOYMON013	g16508	BLASTN	558	1e-37	82
2284	16	700869124H1	SOYMON016	g16508	BLASTN	559	1e-37	83
2285	16	701100219H1	SOYMON028	g2305013	BLASTN	560	1e-37	80
2286	16	700992589H1	SOYMON011	g497899	BLASTN	560	1e-37	80
2287	16	700865642H1	SOYMON016	g16508	BLASTN	562	1e-37	84
2288	16	701068835H1	SOYMON034	g429105	BLASTN	321	1e-36	86
2289	16	700746908H1	SOYMON013	g16508	BLASTN	373	1e-36	81
2290	16	700954719H1	SOYMON022	g16508	BLASTN	475	1e-36	86
2291	16	701042790H1	SOYMON029	g497899	BLASTN	489	1e-36	88
2292	16	700835259H1	SOYMON019	g497899	BLASTN	507	1e-36	88
2293	16	700565816H1	SOYMON002	g16508	BLASTN	519	1e-36	82
2294	16	700895811H1	SOYMON027	g609224	BLASTN	540	1e-36	83
2295	16	700975709H1	SOYMON009	g609224	BLASTN	541	1e-36	84
2296	16	700751645H1	SOYMON014	g16508	BLASTN	542	1e-36	85
2297	16	700567087H1	SOYMON002	g16508	BLASTN	542	1e-36	85
2298	16	700893027H1	SOYMON024	g609224	BLASTN	542	1e-36	85
2299	16	700891185H1	SOYMON024	g609224	BLASTN	543	1e-36	84
2300	16	700851376H1	SOYMON023	g16508	BLASTN	543	1e-36	83
2301	16	700898322H1	SOYMON027	g609224	BLASTN	547	1e-36	85
2302	16	700762881H1	SOYMON015	g16508	BLASTN	323	1e-35	85
2303	16	700960023H1	SOYMON022	g16508	BLASTN	370	1e-35	90
2304	16	701136486H1	SOYMON038	g497899	BLASTN	378	1e-35	85
2305	16	700987688H1	SOYMON009	g497899	BLASTN	387	1e-35	83
2306	16	700836508H1	SOYMON020	g16508	BLASTN	431	1e-35	85
2307	16	700990779H1	SOYMON011	g167961	BLASTN	471	1e-35	86
2308	16	700753019H1	SOYMON014	g497899	BLASTN	476	1e-35	83
2309	16	701138490H1	SOYMON038	g16508	BLASTN	481	1e-35	83
2310	16	700751771H1	SOYMON014	g609224	BLASTN	528	1e-35	84
2311	16	701048324H1	SOYMON032	g609224	BLASTN	529	1e-35	84
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2313	16	700986295H1	SOYMON009	g497899	BLASTN	533	1e-35	86
2314	16	701205693H1	SOYMON035	g16508	BLASTN	535	1e-35	84
2315	16	700865674H1	SOYMON016	g609224	BLASTN	536	1e-35	85
2316	16	700943777H1	SOYMON024	g16508	BLASTN	536	1e-35	84
2317	16	700968207H1	SOYMON035	g609224	BLASTN	536	1e-35	85
2318	16	700898370H1	SOYMON027	g609224	BLASTN	536	1e-35	85
2319	16	700987890H1	SOYMON009	g609224	BLASTN	536	1e-35	85
2320	16	700896016H1	SOYMON027	g609224	BLASTN	536	1e-35	85
2321	16	700554440H1	SOYMON001	g16508	BLASTN	283	1e-34	82
2322	16	700789855H2	SOYMON011	g16508	BLASTN	293	1e-34	91
2323	16	701101393H1	SOYMON028	g16508	BLASTN	311	1e-34	84
2324	16	700946335H1	SOYMON024	g16508	BLASTN	323	1e-34	86
2325	16	701207420H1	SOYMON035	g169664	BLASTN	350	1e-34	88
2326	16	700733034H1	SOYMON010	g16508	BLASTN	384	1e-34	81
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2328	16	700901808H1	SOYMON027	g497899	BLASTN	430	1e-34	81
2329	16	700760954H1	SOYMON015	g609224	BLASTN	514	1e-34	84
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2331	16	701039157H1	SOYMON029	g609224	BLASTN	520	1e-34	76
2332	16	700991056H1	SOYMON011	g166873	BLASTN	520	1e-34	80
2333	16	701097173H1	SOYMON028	g609224	BLASTN	521	1e-34	85

2334	16	700900989H1	SOYMON027	g497899	BLASTN	521	1e-34	86
2335	16	700895584H1	SOYMON027	g609224	BLASTN	521	1e-34	85
2336	16	700726668H1	SOYMON009	g16508	BLASTN	521	1e-34	85
2337	16	701011191H1	SOYMON019	g16508	BLASTN	521	1e-34	85
2338	16	701100827H1	SOYMON028	g16508	BLASTN	522	1e-34	84
2339	16	701156732H1	SOYMON031	g16508	BLASTN	522	1e-34	84
2340	16	700889752H1	SOYMON024	g609224	BLASTN	523	1e-34	84
2341	16	701105873H1	SOYMON036	g16508	BLASTN	524	1e-34	87
2342	16	700958346H1	SOYMON022	g497899	BLASTN	525	1e-34	84
2343	16	700962972H1	SOYMON022	g609224	BLASTN	525	1e-34	84
2344	16	701055918H1	SOYMON032	g16508	BLASTN	526	1e-34	85
2345	16	701123061H1	SOYMON037	g16508	BLASTN	526	1e-34	85
2346	16	701109796H1	SOYMON036	g166873	BLASTN	335	1e-33	86
2347	16	701207350H1	SOYMON035	g497899	BLASTN	349	1e-33	89
2348	16	700835692H1	SOYMON019	g16508	BLASTN	377	1e-33	81
2349	16	700729083H1	SOYMON009	g16508	BLASTN	448	1e-33	85
2350	16	LIB3040-049-Q1-E1-E8	LIB3040	g16508	BLASTN	464	1e-33	80
2351	16	700896567H1	SOYMON027	g497899	BLASTN	502	1e-33	81
2352	16	700668032H1	SOYMON006	g609224	BLASTN	504	1e-33	85
2353	16	700895062H1	SOYMON024	g609224	BLASTN	505	1e-33	83
2354	16	700961178H1	SOYMON022	g16508	BLASTN	506	1e-33	84
2355	16	701202590H1	SOYMON035	g497899	BLASTN	508	1e-33	86
2356	16	700754960H1	SOYMON014	g16508	BLASTN	512	1e-33	83
2357	16	700748584H1	SOYMON013	g16508	BLASTN	512	1e-33	84
2358	16	700791669H1	SOYMON011	g609224	BLASTN	512	1e-33	84
2359	16	701210405H1	SOYMON035	g16508	BLASTN	526	1e-33	83
2360	16	LIB3040-054-Q1-E1-D8	LIB3040	g16508	BLASTN	531	1e-33	80
2361	16	700791176H1	SOYMON011	g497899	BLASTN	311	1e-32	87
2362	16	LIB3049-029-Q1-E1-C5	LIB3049	g167961	BLASTN	368	1e-32	73
2363	16	700893458H1	SOYMON024	g497899	BLASTN	380	1e-32	87
2364	16	700829728H1	SOYMON019	g497899	BLASTN	490	1e-32	88
2365	16	700833137H1	SOYMON019	g609224	BLASTN	490	1e-32	85
2366	16	701204592H2	SOYMON035	g609224	BLASTN	493	1e-32	86
2367	16	700834821H1	SOYMON019	g609224	BLASTN	494	1e-32	78
2368	16	700757167H1	SOYMON015	g16508	BLASTN	494	1e-32	85
2369	16	701203730H2	SOYMON035	g16508	BLASTN	496	1e-32	82
2370	16	701044359H1	SOYMON032	g609224	BLASTN	497	1e-32	84
2371	16	701010095H2	SOYMON019	g16508	BLASTN	498	1e-32	86
2372	16	701100059H2	SOYMON028	g16508	BLASTN	499	1e-32	80
2373	16	700844256H1	SOYMON021	g16508	BLASTN	500	1e-32	80
2374	16	700968296H1	SOYMON035	g497899	BLASTN	501	1e-32	85
2375	16	701110404H1	SOYMON036	g16508	BLASTN	333	1e-31	84
2376	16	700869025H1	SOYMON016	g497899	BLASTN	340	1e-31	86
2377	16	700556245H1	SOYMON001	g609224	BLASTN	355	1e-31	84
2378	16	700566166H1	SOYMON002	g726031	BLASTN	402	1e-31	77
2379	16	700941481H1	SOYMON024	g497899	BLASTN	423	1e-31	79
2380	16	700896940H1	SOYMON027	g169664	BLASTN	477	1e-31	85
2381	16	701145429H1	SOYMON031	g609224	BLASTN	478	1e-31	85
2382	16	700946496H1	SOYMON024	g609224	BLASTN	488	1e-31	86
2383	16	700902149H1	SOYMON027	g1655577	BLASTN	488	1e-31	87
2384	16	700653072H1	SOYMON003	g497899	BLASTN	489	1e-31	88
2385	16	701000261H1	SOYMON018	g16508	BLASTN	495	1e-31	78
2386	16	701039358H1	SOYMON029	g609224	BLASTN	320	1e-30	85
2387	16	701156951H1	SOYMON031	g609224	BLASTN	467	1e-30	86

2388	16	701040880H1	SOYMON029	g497899	BLASTN	477	1e-30	89
2389	16	700979632H2	SOYMON009	g609224	BLASTN	490	1e-30	83
2390	16	700983678H1	SOYMON009	g16508	BLASTN	192	1e-29	86
2391	16	700893644H1	SOYMON024	g609224	BLASTN	320	1e-29	85
2392	16	700962437H1	SOYMON022	g609224	BLASTN	364	1e-29	83
2393	16	701150545H1	SOYMON031	g16508	BLASTN	457	1e-29	86
2394	16	700555403H1	SOYMON001	g16508	BLASTN	458	1e-29	90
2395	16	700794307H1	SOYMON017	g16508	BLASTN	460	1e-29	87
2396	16	701137971H1	SOYMON038	g16508	BLASTN	460	1e-29	84
2397	16	700893774H1	SOYMON024	g609224	BLASTN	461	1e-29	86
2398	16	700565272H1	SOYMON002	g609224	BLASTN	464	1e-29	74
2399	16	700667161H1	SOYMON006	g16508	BLASTN	468	1e-29	75
2400	16	700752952H1	SOYMON014	g609224	BLASTN	279	1e-28	78
2401	16	700901534H1	SOYMON027	g2305013	BLASTN	288	1e-28	85
2402	16	701006617H1	SOYMON019	g497899	BLASTN	358	1e-28	79
2403	16	700753459H1	SOYMON014	g609224	BLASTN	446	1e-28	81
2404	16	701157089H1	SOYMON031	g609224	BLASTN	448	1e-28	84
2405	16	700831271H1	SOYMON019	g16508	BLASTN	453	1e-28	87
2406	16	701207325H1	SOYMON035	g167961	BLASTN	251	1e-27	81
2407	16	700566121H1	SOYMON002	g16508	BLASTN	259	1e-27	75
2408	16	701137878H1	SOYMON038	g609224	BLASTN	280	1e-27	85
2409	16	700763957H1	SOYMON019	g497899	BLASTN	430	1e-27	88
2410	16	701015858H1	SOYMON038	g609224	BLASTN	431	1e-27	86
2411	16	700742849H1	SOYMON012	g497899	BLASTN	434	1e-27	88
2412	16	701102648H1	SOYMON028	g16508	BLASTN	436	1e-27	89
2413	16	700940955H1	SOYMON024	g609556	BLASTN	438	1e-27	85
2414	16	700889805H1	SOYMON024	g16508	BLASTN	441	1e-27	89
2415	16	701061930H1	SOYMON033	g16508	BLASTN	315	1e-26	77
2416	16	700901375H1	SOYMON027	g609224	BLASTN	323	1e-26	84
2417	16	700989056H1	SOYMON011	g609224	BLASTN	345	1e-26	82
2418	16	700665988H1	SOYMON005	g960356	BLASTN	420	1e-26	79
2419	16	700893352H1	SOYMON024	g497899	BLASTN	428	1e-26	87
2420	16	701122945H1	SOYMON037	g16508	BLASTN	269	1e-25	76
2421	16	700992929H1	SOYMON011	g16508	BLASTN	328	1e-25	78
2422	16	700557564H1	SOYMON001	g609224	BLASTN	413	1e-25	85
2423	16	701102620H1	SOYMON028	g609224	BLASTN	414	1e-25	82
2424	16	701062321H1	SOYMON033	g497899	BLASTN	267	1e-24	87
2425	16	701062258H1	SOYMON033	g450548	BLASTN	391	1e-24	83
2426	16	701145703H1	SOYMON031	g609224	BLASTN	411	1e-24	85
2427	16	701144943H1	SOYMON031	g450548	BLASTN	421	1e-24	85
2428	16	701134190H1	SOYMON038	g2305013	BLASTN	268	1e-23	87
2429	16	701132116H1	SOYMON038	g450548	BLASTN	385	1e-23	85
2430	16	701144522H1	SOYMON031	g450548	BLASTN	406	1e-23	85
2431	16	700738002H1	SOYMON012	g1655578	BLASTX	205	1e-21	84
2432	16	700756686H1	SOYMON014	g16508	BLASTN	235	1e-21	76
2433	16	700649058H1	SOYMON003	g16508	BLASTN	360	1e-21	85
2434	16	701152050H1	SOYMON031	g497899	BLASTN	233	1e-20	87
2435	16	701156991H1	SOYMON031	g497900	BLASTX	146	1e-19	98
2436	16	700761949H1	SOYMON015	g497900	BLASTX	173	1e-19	98
2437	16	700988163H1	SOYMON009	g609224	BLASTN	243	1e-19	81
2438	16	700666050H1	SOYMON005	g960357	BLASTX	183	1e-18	89
2439	16	700896896H1	SOYMON027	g497900	BLASTX	95	1e-17	82
2440	16	701100249H1	SOYMON028	g497900	BLASTX	104	1e-17	79
2441	16	701061719H1	SOYMON033	g609224	BLASTN	310	1e-17	67

2442	16	700674836H1	SOYMON007	g16508	BLASTN	331	1e-17	84
2443	16	700908822H1	SOYMON022	g169665	BLASTX	171	1e-16	100
2444	16	700898670H1	SOYMON027	g166872	BLASTX	172	1e-16	93
2445	16	700650967H1	SOYMON003	g16845	BLASTX	147	1e-15	100
2446	16	701132185H1	SOYMON038	g16845	BLASTX	151	1e-15	100
2447	16	700648929H1	SOYMON003	g1033190	BLASTX	156	1e-14	96
2448	16	700960809H1	SOYMON022	g16508	BLASTN	168	1e-14	88
2449	16	701101753H1	SOYMON028	g16845	BLASTX	141	1e-13	88
2450	16	700742010H1	SOYMON012	g609224	BLASTN	273	1e-13	88
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2452	16	700979242H1	SOYMON009	g609224	BLASTN	165	1e-12	81
2453	16	700740838H1	SOYMON012	g609224	BLASTN	253	1e-12	88
2454	16	700832383H1	SOYMON019	g16845	BLASTX	82	1e-11	71
2455	16	700830938H1	SOYMON019	g16845	BLASTX	92	1e-11	90
2456	16	700564686H1	SOYMON002	g1655578	BLASTX	95	1e-11	97
2457	16	700753178H1	SOYMON014	g16845	BLASTX	107	1e-10	78
2458	16	700563834H1	SOYMON002	g497900	BLASTX	124	1e-10	100
2459	16	700897739H1	SOYMON027	g16845	BLASTX	125	1e-10	86
2460	16	700725722H1	SOYMON009	g609225	BLASTX	125	1e-10	92
2461	16	701210357H1	SOYMON035	g497900	BLASTX	128	1e-10	100
2462	16	700833654H1	SOYMON019	g609224	BLASTN	243	1e-10	87
2463	16	701010334H1	SOYMON019	g609224	BLASTN	258	1e-10	86
2464	16	700567622H1	SOYMON002	g166874	BLASTX	93	1e-9	100
2465	16	700647933H1	SOYMON003	g609225	BLASTX	120	1e-9	81
2466	16	700742255H1	SOYMON012	g429107	BLASTN	160	1e-9	86
2467	16	700981506H1	SOYMON009	g609224	BLASTN	248	1e-9	88
2468	16	700962044H1	SOYMON022	g497900	BLASTX	113	1e-8	100
2469	16	701039604H1	SOYMON029	g497900	BLASTX	113	1e-8	100
2470	16	701009941H2	SOYMON019	g609225	BLASTX	114	1e-8	70
2471	16	700976956H1	SOYMON009	g609225	BLASTX	118	1e-8	96
2472	18138	701120722H1	SOYMON037	g169664	BLASTN	514	1e-33	92
2473	18138	700946044H1	SOYMON024	g169664	BLASTN	437	1e-26	90
2474	18138	700664443H1	SOYMON005	g17262	BLASTX	162	1e-16	88
2475	18138	700665162H1	SOYMON005	g17262	BLASTX	162	1e-16	88
2476	18138	701143667H1	SOYMON038	g16961	BLASTX	123	1e-11	95
2477	18138	701099150H1	SOYMON028	g16961	BLASTX	112	1e-9	90
2478	27686	700909141H1	SOYMON022	g726027	BLASTN	723	1e-51	84
2479	27686	701145458H1	SOYMON031	g726027	BLASTN	694	1e-49	86

ADENOSYLMETHIONINE DECARBOXYLASE (EC 4.1.1.50)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
430	-700151703	700151703H1	SATMON007	g1532072	BLASTN	717	1e-50	91
431	-700165608	700165608H1	SATMON013	g1532072	BLASTN	605	1e-41	75
432	-700166868	700166868H1	SATMON013	g1532072	BLASTN	593	1e-40	90
433	-700242148	700242148H1	SATMON010	g1532048	BLASTX	142	1e-12	88
434	-700354923	700354923H1	SATMON024	g1532072	BLASTN	887	1e-74	89
435	-700422279	700422279H1	SATMONN01	g1532072	BLASTN	623	1e-61	88
436	-700455682	700455682H1	SATMON029	g1532047	BLASTN	681	1e-47	77
437	-700477645	700477645H1	SATMON025	g1532048	BLASTX	90	1e-12	71
438	-700550509	700550509H1	SATMON022	g1532047	BLASTN	399	1e-41	80
439	-700572502	700572502H1	SATMON030	g1532048	BLASTX	93	1e-20	70
440	-700618258	700618258H1	SATMON033	g1532072	BLASTN	368	1e-67	92

441	-701165456	701165456H1	SATMONN04	g1532072	BLASTN	436	1e-26	71
442	-L30622289	LIB3062-004-Q1-K1-E10	LIB3062	g1532047	BLASTN	406	1e-26	82
443	-L30623185	LIB3062-026-Q1-K1-E2	LIB3062	g1532072	BLASTN	338	1e-19	62
444	-L30625601	LIB3062-033-Q1-K1-A9	LIB3062	g1403043	BLASTN	758	1e-54	68
445	-L30661842	LIB3066-009-Q1-K1-E4	LIB3066	g1532072	BLASTN	398	1e-22	86
446	-L30681932	LIB3068-020-Q1-K1-C8	LIB3068	g1532072	BLASTN	205	1e-28	89
447	-L30687176	LIB3068-059-Q1-K1-C1	LIB3068	g1532072	BLASTN	216	1e-20	87
448	-L30692075	LIB3069-004-Q1-K1-G11	LIB3069	g1532072	BLASTN	619	1e-40	90
449	-L30783194	LIB3078-052-Q1-K1-E4	LIB3078	g1532072	BLASTN	447	1e-40	76
450	-L30792336	LIB3079-021-Q1-K1-E12	LIB3079	g1532072	BLASTN	217	1e-13	80
451	1471	700106762H1	SATMON010	g1532072	BLASTN	323	1e-59	84
452	1471	701163744H1	SATMONN04	g1532072	BLASTN	220	1e-41	86
453	1471	LIB3059-023-Q1-K1-A11	LIB3059	g1532072	BLASTN	340	1e-39	80
454	1471	700574530H1	SATMON030	g1532072	BLASTN	247	1e-34	82
455	1471	LIB83-003-Q1-E1-G1	LIB83	g1532072	BLASTN	239	1e-24	80
456	1471	701161147H1	SATMONN04	g1532072	BLASTN	239	1e-16	80
457	1471	700477336H1	SATMON025	g1532072	BLASTN	235	1e-10	91
458	16729	700169502H1	SATMON013	g1403043	BLASTN	456	1e-28	65
459	16729	700088201H1	SATMON011	g1532048	BLASTX	123	1e-22	60
460	16729	700165440H1	SATMON013	g1532048	BLASTX	162	1e-15	65
461	16866	700072905H1	SATMON007	g1532047	BLASTN	368	1e-19	71
462	16866	700350558H1	SATMON023	g1532047	BLASTN	301	1e-14	70
463	16866	700088370H1	SATMON011	g1532047	BLASTN	306	1e-14	70
464	2324	LIB3066-042-Q1-K1-H2	LIB3066	g1403043	BLASTN	1344	1e-103	79
465	2324	LIB3059-011-Q1-K1-B2	LIB3059	g1403043	BLASTN	1351	1e-103	78
466	2324	LIB3059-042-Q1-K1-B7	LIB3059	g1403043	BLASTN	1230	1e-100	80
467	2324	LIB3062-036-Q1-K1-F2	LIB3062	g1532072	BLASTN	1097	1e-91	80
468	2324	LIB3069-020-Q1-K1-A5	LIB3069	g1403043	BLASTN	925	1e-82	80
469	2324	LIB189-023-Q1-E1-H3	LIB189	g1532072	BLASTN	1085	1e-81	78
470	2324	700266088H1	SATMON017	g1532047	BLASTN	998	1e-74	80
471	2324	LIB3062-026-Q1-K1-E6	LIB3062	g1532047	BLASTN	982	1e-72	77
472	2324	700083335H1	SATMON011	g1403043	BLASTN	960	1e-71	79
473	2324	LIB189-015-Q1-E1-D3	LIB189	g1532047	BLASTN	970	1e-71	77
474	2324	LIB3079-013-Q1-K1-B3	LIB3079	g1403043	BLASTN	953	1e-70	81
475	2324	700262129H1	SATMON017	g1403043	BLASTN	936	1e-69	79
476	2324	700265667H1	SATMON017	g1403043	BLASTN	694	1e-66	81
477	2324	700807255H1	SATMON036	g1532072	BLASTN	850	1e-62	81
478	2324	700196129H1	SATMON014	g1403043	BLASTN	853	1e-62	83
479	2324	LIB3066-042-Q1-K1-H1	LIB3066	g1403043	BLASTN	842	1e-61	80
480	2324	700458321H1	SATMON029	g1403043	BLASTN	501	1e-58	80
481	2324	700197643H1	SATMON014	g1403043	BLASTN	461	1e-57	80
482	2324	LIB143-029-Q1-E1-H3	LIB143	g1532072	BLASTN	786	1e-56	78
483	2324	700197842H1	SATMON014	g1403043	BLASTN	688	1e-48	85
484	2324	700196807H1	SATMON014	g1532072	BLASTN	678	1e-47	75
485	2324	700263712H1	SATMON017	g1532072	BLASTN	403	1e-45	75
486	2324	LIB143-006-Q1-E1-F9	LIB143	g1403043	BLASTN	324	1e-42	83
487	2324	700267496H1	SATMON017	g1532047	BLASTN	594	1e-40	83
488	2324	700172551H1	SATMON013	g1532072	BLASTN	546	1e-36	74
489	2324	700211893H1	SATMON016	g1532047	BLASTN	542	1e-35	82
490	2324	700264065H1	SATMON017	g1532047	BLASTN	515	1e-34	83
491	2324	700465305H1	SATMON025	g1532073	BLASTX	148	1e-26	61
492	2324	700455052H1	SATMON029	g1403043	BLASTN	274	1e-26	79
493	2324	700473305H1	SATMON025	g1403043	BLASTN	446	1e-26	74
494	2324	700475851H1	SATMON025	g1532047	BLASTN	333	1e-18	78

495	2324	700263111H1	SATMON017	g1532073	BLASTX	165	1e-15	72
496	2324	700465705H1	SATMON025	g1403044	BLASTX	78	1e-8	76
497	3185	700264424H1	SATMON017	g1403043	BLASTN	469	1e-41	81
498	3185	LIB143-007-Q1-E1-H2	LIB143	g1403043	BLASTN	464	1e-36	80
499	3185	700262548H1	SATMON017	g1532047	BLASTN	368	1e-34	80
500	3185	700263845H1	SATMON017	g1403043	BLASTN	455	1e-34	81
501	3185	LIB3068-025-Q1-K1-G10	LIB3068	g1403043	BLASTN	288	1e-32	81
502	3185	700264569H1	SATMON017	g1403043	BLASTN	469	1e-32	82
503	3185	700243571H1	SATMON010	g1532047	BLASTN	461	1e-28	77
504	3185	700265453H1	SATMON017	g1532047	BLASTN	269	1e-27	84
505	3185	700267779H1	SATMON017	g1532047	BLASTN	257	1e-26	85
506	3185	700382373H1	SATMON024	g1532047	BLASTN	255	1e-24	86
507	3185	700258727H1	SATMON017	g1532047	BLASTN	255	1e-24	84
508	3185	LIB3069-018-Q1-K1-F5	LIB3069	g1532047	BLASTN	255	1e-21	77
509	3185	LIB3066-038-Q1-K1-C2	LIB3066	g1403043	BLASTN	242	1e-17	68
510	3185	700334654H1	SATMON019	g1532047	BLASTN	248	1e-17	90
511	3185	700262830H1	SATMON017	g1403043	BLASTN	276	1e-12	78
512	3185	700264727H1	SATMON017	g1532047	BLASTN	255	1e-10	92
513	3185	700238407H1	SATMON010	g1532047	BLASTN	255	1e-10	92
514	3185	700441952H1	SATMON026	g1532047	BLASTN	255	1e-10	92
515	3185	700268188H1	SATMON017	g1532047	BLASTN	255	1e-10	92
516	3185	700801627H1	SATMON036	g1532047	BLASTN	255	1e-10	92
517	3185	700261609H1	SATMON017	g1532047	BLASTN	255	1e-10	92
518	3185	700258510H1	SATMON017	g1532047	BLASTN	255	1e-10	92
519	3185	700258779H1	SATMON017	g1532047	BLASTN	255	1e-10	92
520	3185	700256838H1	SATMON017	g1532047	BLASTN	255	1e-10	92
521	3185	700263361H1	SATMON017	g1532047	BLASTN	255	1e-10	92
522	3185	700257102H1	SATMON017	g1532047	BLASTN	255	1e-10	92
523	3185	700239452H1	SATMON010	g1532047	BLASTN	245	1e-9	91
524	3185	700262045H1	SATMON017	g1532047	BLASTN	250	1e-9	90
525	8	LIB3066-027-Q1-K1-C6	LIB3066	g1532072	BLASTN	1414	1e-181	97
526	8	LIB3066-048-Q1-K1-C9	LIB3066	g1532072	BLASTN	1715	1e-173	98
527	8	LIB3066-007-Q1-K1-B4	LIB3066	g1532072	BLASTN	2146	1e-170	97
528	8	LIB3066-019-Q1-K1-B9	LIB3066	g1532072	BLASTN	1884	1e-168	98
529	8	LIB148-038-Q1-E1-A3	LIB148	g1532072	BLASTN	1585	1e-164	99
530	8	LIB3068-002-Q1-K1-H5	LIB3068	g1532072	BLASTN	2051	1e-162	98
531	8	LIB3067-035-Q1-K1-E9	LIB3067	g1532072	BLASTN	1594	1e-161	98
532	8	LIB189-020-Q1-E1-E8	LIB189	g1532072	BLASTN	1501	1e-160	98
533	8	LIB3078-057-Q1-K1-C12	LIB3078	g1532072	BLASTN	2026	1e-160	96
534	8	LIB3066-053-Q1-K1-A8	LIB3066	g1532072	BLASTN	1713	1e-159	93
535	8	LIB189-006-Q1-E1-C6	LIB189	g1532072	BLASTN	1676	1e-157	98
536	8	LIB3078-054-Q1-K1-E4	LIB3078	g1532072	BLASTN	1997	1e-157	95
537	8	LIB3069-028-Q1-K1-A11	LIB3069	g1532072	BLASTN	1980	1e-156	96
538	8	LIB3069-045-Q1-K1-H7	LIB3069	g1532072	BLASTN	1989	1e-156	98
539	8	LIB148-025-Q1-E1-F7	LIB148	g1532072	BLASTN	1954	1e-154	97
540	8	LIB189-014-Q1-E1-F11	LIB189	g1532072	BLASTN	1925	1e-151	94
541	8	LIB3060-014-Q1-K1-C5	LIB3060	g1532072	BLASTN	1780	1e-150	96
542	8	LIB148-057-Q1-E1-C2	LIB148	g1532072	BLASTN	1000	1e-149	98
543	8	LIB148-015-Q1-E1-F2	LIB148	g1532072	BLASTN	1886	1e-148	94
544	8	LIB3066-045-Q1-K1-A2	LIB3066	g1532072	BLASTN	914	1e-146	91
545	8	LIB148-064-Q1-E1-A3	LIB148	g1532072	BLASTN	1660	1e-146	93
546	8	LIB3061-047-Q1-K1-G1	LIB3061	g1532072	BLASTN	1867	1e-146	92
547	8	LIB148-040-Q1-E1-F1	LIB148	g1532072	BLASTN	1706	1e-145	94
548	8	LIB3069-012-Q1-K1-B6	LIB3069	g1532072	BLASTN	1630	1e-144	87

549	8	LIB3068-009-Q1-K1-A8	LIB3068	g1532072	BLASTN	1802	1e-141	97
550	8	LIB148-004-Q1-E1-D2	LIB148	g1532072	BLASTN	818	1e-139	91
551	8	LIB3068-002-Q1-K1-A1	LIB3068	g1532072	BLASTN	944	1e-137	95
552	8	LIB3067-035-Q1-K1-H9	LIB3067	g1532072	BLASTN	1701	1e-137	98
553	8	LIB3068-033-Q1-K1-G12	LIB3068	g1532072	BLASTN	1093	1e-130	89
554	8	700572229H1	SATMON030	g1532072	BLASTN	1135	1e-128	99
555	8	LIB3078-052-Q1-K1-E1	LIB3078	g1532072	BLASTN	1235	1e-127	85
556	8	700572579H1	SATMON030	g1532072	BLASTN	1625	1e-126	98
557	8	LIB3066-025-Q1-K1-F2	LIB3066	g1532072	BLASTN	1625	1e-126	100
558	8	LIB3068-048-Q1-K1-F9	LIB3068	g1532072	BLASTN	1538	1e-125	98
559	8	700098413H1	SATMON009	g1532072	BLASTN	1595	1e-124	100
560	8	700573235H1	SATMON030	g1532072	BLASTN	1513	1e-123	98
561	8	700090946H1	SATMON011	g1532072	BLASTN	1585	1e-123	100
562	8	700092465H1	SATMON008	g1532072	BLASTN	1541	1e-122	98
563	8	700074625H1	SATMON007	g1532072	BLASTN	1570	1e-122	100
564	8	LIB3059-007-Q1-K1-C10	LIB3059	g1532072	BLASTN	1401	1e-119	94
565	8	700072828H1	SATMON007	g1532072	BLASTN	1540	1e-119	100
566	8	700619106H1	SATMON034	g1532072	BLASTN	833	1e-118	97
567	8	700074853H1	SATMON007	g1532072	BLASTN	1515	1e-117	100
568	8	700201293H1	SATMON003	g1532072	BLASTN	1517	1e-117	97
569	8	700075896H1	SATMON007	g1532072	BLASTN	1006	1e-115	99
570	8	LIB3059-052-Q1-K1-A1	LIB3059	g1532072	BLASTN	1241	1e-115	92
571	8	700091576H1	SATMON011	g1532072	BLASTN	943	1e-114	98
572	8	LIB3078-015-Q1-K1-D7	LIB3078	g1532072	BLASTN	1218	1e-114	87
573	8	700074733H1	SATMON007	g1532072	BLASTN	1475	1e-114	100
574	8	700381421H1	SATMON023	g1532072	BLASTN	1475	1e-114	100
575	8	700085594H1	SATMON011	g1532072	BLASTN	1477	1e-114	99
576	8	700095883H1	SATMON008	g1532072	BLASTN	1477	1e-114	99
577	8	700338237H1	SATMON020	g1532072	BLASTN	1478	1e-114	99
578	8	700549813H1	SATMON022	g1532072	BLASTN	1481	1e-114	99
579	8	700097935H1	SATMON009	g1532072	BLASTN	1484	1e-114	98
580	8	700572978H1	SATMON030	g1532072	BLASTN	865	1e-113	96
581	8	700196464H1	SATMON014	g1532072	BLASTN	1044	1e-113	92
582	8	700381412H1	SATMON023	g1532072	BLASTN	1168	1e-113	98
583	8	700027839H1	SATMON003	g1532072	BLASTN	1472	1e-113	99
584	8	700623344H1	SATMON034	g1532072	BLASTN	1085	1e-112	94
585	8	700025858H1	SATMON003	g1532072	BLASTN	1455	1e-112	100
586	8	LIB3059-017-Q1-K1-H5	LIB3059	g1532072	BLASTN	1459	1e-112	97
587	8	700475019H1	SATMON025	g1532072	BLASTN	1390	1e-111	100
588	8	700338357H1	SATMON020	g1532072	BLASTN	1440	1e-111	100
589	8	700256796H1	SATMON017	g1532072	BLASTN	1400	1e-110	100
590	8	700071692H1	SATMON007	g1532072	BLASTN	1430	1e-110	98
591	8	700339073H1	SATMON020	g1532072	BLASTN	1357	1e-109	95
592	8	700106916H1	SATMON010	g1532072	BLASTN	1415	1e-109	93
593	8	700468234H1	SATMON025	g1532072	BLASTN	1420	1e-109	100
594	8	700214482H1	SATMON016	g1532072	BLASTN	1425	1e-109	100
595	8	700466437H1	SATMON025	g1532072	BLASTN	1403	1e-108	98
596	8	700205576H1	SATMON003	g1532072	BLASTN	1404	1e-108	98
597	8	700043455H1	SATMON004	g1532072	BLASTN	1405	1e-108	100
598	8	700348439H1	SATMON023	g1532072	BLASTN	1407	1e-108	99
599	8	700093131H1	SATMON008	g1532072	BLASTN	755	1e-107	100
600	8	700571851H1	SATMON030	g1532072	BLASTN	1302	1e-107	99
601	8	700088142H1	SATMON011	g1532072	BLASTN	1392	1e-107	99
602	8	700085934H1	SATMON011	g1532072	BLASTN	1397	1e-107	98

603	8	700028465H1	SATMON003	g1532072	BLASTN	1258	1e-106	98
604	8	700236818H1	SATMON010	g1532072	BLASTN	1380	1e-106	100
605	8	700583691H1	SATMON031	g1532072	BLASTN	1382	1e-106	99
606	8	700095585H1	SATMON008	g1532072	BLASTN	1383	1e-106	94
607	8	700105140H1	SATMON010	g1532072	BLASTN	1375	1e-105	100
608	8	700338486H1	SATMON020	g1532072	BLASTN	893	1e-104	98
609	8	700214178H1	SATMON016	g1532072	BLASTN	1361	1e-104	99
610	8	700090613H1	SATMON011	g1532072	BLASTN	699	1e-103	99
611	8	700576189H1	SATMON030	g1532072	BLASTN	1254	1e-103	93
612	8	700475734H1	SATMON025	g1532072	BLASTN	1256	1e-103	99
613	8	700028218H1	SATMON003	g1532072	BLASTN	1275	1e-103	100
614	8	700088644H1	SATMON011	g1532072	BLASTN	1351	1e-103	98
615	8	700378768H1	SATMON020	g1532072	BLASTN	1067	1e-102	98
616	8	LIB3067-035-Q1-K1-H10	LIB3067	g1532072	BLASTN	1233	1e-102	95
617	8	700043466H1	SATMON004	g1532072	BLASTN	1331	1e-102	97
618	8	LIB148-057-Q1-E1-C3	LIB148	g1532072	BLASTN	1283	1e-101	92
619	8	700217574H1	SATMON016	g1532072	BLASTN	1320	1e-101	100
620	8	700042332H1	SATMON004	g1532072	BLASTN	1321	1e-101	97
621	8	700440926H1	SATMON026	g1532072	BLASTN	1323	1e-101	99
622	8	700552284H1	SATMON022	g1532072	BLASTN	1329	1e-101	97
623	8	700216613H1	SATMON016	g1532072	BLASTN	689	1e-100	97
624	8	LIB3067-054-Q1-K1-F6	LIB3067	g1532072	BLASTN	1030	1e-100	94
625	8	LIB3060-038-Q1-K1-B9	LIB3060	g1532072	BLASTN	1108	1e-100	92
626	8	700218755H1	SATMON011	g1532072	BLASTN	1116	1e-100	99
627	8	700338042H1	SATMON020	g1532072	BLASTN	1270	1e-100	99
628	8	700268009H1	SATMON017	g1532072	BLASTN	1307	1e-100	92
629	8	700578491H1	SATMON031	g1532072	BLASTN	1309	1e-100	97
630	8	700030037H1	SATMON003	g1532072	BLASTN	1310	1e-100	97
631	8	700221406H1	SATMON011	g1532072	BLASTN	1315	1e-100	100
632	8	700157357H1	SATMON012	g1532072	BLASTN	1315	1e-100	98
633	8	700476926H1	SATMON025	g1532072	BLASTN	680	1e-99	97
634	8	700475068H1	SATMON025	g1532072	BLASTN	792	1e-99	98
635	8	700469741H1	SATMON025	g1532072	BLASTN	831	1e-99	99
636	8	700082377H1	SATMON011	g1532072	BLASTN	1081	1e-99	99
637	8	700217143H1	SATMON016	g1532072	BLASTN	1208	1e-99	98
638	8	700339433H1	SATMON020	g1532072	BLASTN	1295	1e-99	100
639	8	700196627H1	SATMON014	g1532072	BLASTN	1295	1e-99	100
640	8	700259354H1	SATMON017	g1532072	BLASTN	1305	1e-99	92
641	8	700610842H1	SATMON022	g1532072	BLASTN	843	1e-98	97
642	8	700218059H1	SATMON016	g1532072	BLASTN	1036	1e-98	99
643	8	700421727H1	SATMONN01	g1532072	BLASTN	1172	1e-98	97
644	8	700158660H1	SATMON012	g1532072	BLASTN	1285	1e-98	100
645	8	700806856H1	SATMON036	g1532072	BLASTN	1194	1e-97	97
646	8	700583512H1	SATMON031	g1532072	BLASTN	1219	1e-97	97
647	8	700025502H1	SATMON004	g1532072	BLASTN	1276	1e-97	99
648	8	700159562H1	SATMON012	g1532072	BLASTN	1277	1e-97	99
649	8	700223731H1	SATMON011	g1532072	BLASTN	1277	1e-97	99
650	8	700156494H1	SATMON012	g1532072	BLASTN	1280	1e-97	100
651	8	700160533H1	SATMON012	g1532072	BLASTN	1281	1e-97	97
652	8	LIB3066-055-Q1-K1-D8	LIB3066	g1532072	BLASTN	672	1e-96	99
653	8	700405152H1	SATMON028	g1532072	BLASTN	828	1e-96	98
654	8	LIB148-040-Q1-E1-A8	LIB148	g1532072	BLASTN	1268	1e-96	83
655	8	700438437H1	SATMON026	g1532072	BLASTN	1246	1e-95	99
656	8	700267245H1	SATMON017	g1532072	BLASTN	1248	1e-95	93

657	8	700156129H2	SATMON007	g1532072	BLASTN	1250	1e-95	100
658	8	700203246H1	SATMON003	g1532072	BLASTN	1250	1e-95	100
659	8	700168339H1	SATMON013	g1532072	BLASTN	915	1e-94	98
660	8	LIB3067-036-Q1-K1-F9	LIB3067	g1532072	BLASTN	1110	1e-94	95
661	8	700193769H1	SATMON014	g1532072	BLASTN	1240	1e-94	100
662	8	700094014H1	SATMON008	g1532072	BLASTN	1241	1e-94	91
663	8	700020311H1	SATMON001	g1532072	BLASTN	1228	1e-93	99
664	8	700195083H1	SATMON014	g1532072	BLASTN	1229	1e-93	98
665	8	700193901H1	SATMON014	g1532072	BLASTN	1229	1e-93	98
666	8	700194639H1	SATMON014	g1532072	BLASTN	1231	1e-93	99
667	8	700350117H1	SATMON023	g1532072	BLASTN	1143	1e-92	99
668	8	700239867H1	SATMON010	g1532072	BLASTN	1210	1e-92	98
669	8	LIB3069-028-Q1-K1-D1	LIB3069	g1532072	BLASTN	1217	1e-92	94
670	8	700355756H1	SATMON024	g1532072	BLASTN	618	1e-91	96
671	8	700265492H1	SATMON017	g1532072	BLASTN	620	1e-91	94
672	8	700579021H1	SATMON031	g1532072	BLASTN	656	1e-91	96
673	8	LIB3079-015-Q1-K1-B11	LIB3079	g1532072	BLASTN	1008	1e-91	83
674	8	700572888H2	SATMON030	g1532072	BLASTN	1159	1e-91	99
675	8	701183990H1	SATMONN06	g1532072	BLASTN	1198	1e-91	93
676	8	700085757H1	SATMON011	g1532072	BLASTN	1200	1e-91	100
677	8	700162756H1	SATMON013	g1532072	BLASTN	1208	1e-91	99
678	8	700193076H1	SATMON014	g1532072	BLASTN	1208	1e-91	99
679	8	700224378H1	SATMON011	g1532072	BLASTN	696	1e-90	99
680	8	700218773H1	SATMON011	g1532072	BLASTN	1187	1e-90	93
681	8	700159338H1	SATMON012	g1532072	BLASTN	1193	1e-90	97
682	8	700169217H1	SATMON013	g1532072	BLASTN	1196	1e-90	99
683	8	700551548H1	SATMON022	g1532072	BLASTN	839	1e-89	96
684	8	700197059H1	SATMON014	g1532072	BLASTN	1025	1e-89	95
685	8	700469834H1	SATMON025	g1532072	BLASTN	650	1e-88	97
686	8	700569778H1	SATMON030	g1532072	BLASTN	682	1e-88	91
687	8	700194845H1	SATMON014	g1532072	BLASTN	1171	1e-88	99
688	8	700445861H1	SATMON027	g1532072	BLASTN	536	1e-87	99
689	8	700100536H1	SATMON009	g1532072	BLASTN	666	1e-87	94
690	8	701163801H1	SATMONN04	g1532072	BLASTN	805	1e-87	94
691	8	LIB3060-035-Q1-K1-E3	LIB3060	g1532072	BLASTN	922	1e-87	95
692	8	700170872H1	SATMON013	g1532072	BLASTN	943	1e-87	97
693	8	700241270H1	SATMON010	g1532072	BLASTN	1066	1e-86	96
694	8	700457981H1	SATMON029	g1532072	BLASTN	1140	1e-86	92
695	8	700158339H1	SATMON012	g1532072	BLASTN	1142	1e-86	99
696	8	700019442H1	SATMON001	g1532072	BLASTN	1145	1e-86	100
697	8	700149885H1	SATMON007	g1532072	BLASTN	1133	1e-85	99
698	8	700018255H1	SATMON001	g1532072	BLASTN	1135	1e-85	100
699	8	700244026H1	SATMON010	g1532072	BLASTN	1044	1e-84	89
700	8	LIB3060-027-Q1-K1-B2	LIB3060	g1532072	BLASTN	1117	1e-84	99
701	8	700170960H1	SATMON013	g1532072	BLASTN	696	1e-83	99
702	8	700267487H1	SATMON017	g1532072	BLASTN	1061	1e-83	92
703	8	700152754H1	SATMON007	g1532072	BLASTN	1107	1e-83	99
704	8	700378260H1	SATMON019	g1532072	BLASTN	1112	1e-83	92
705	8	700455089H1	SATMON029	g1532072	BLASTN	583	1e-82	97
706	8	700167322H1	SATMON013	g1532072	BLASTN	1080	1e-81	98
707	8	LIB3060-035-Q1-K1-H1	LIB3060	g1532072	BLASTN	782	1e-80	94
708	8	700204067H1	SATMON003	g1532072	BLASTN	1011	1e-80	98
709	8	700049271H1	SATMON003	g1532072	BLASTN	627	1e-79	93
710	8	700442065H1	SATMON026	g1532072	BLASTN	1043	1e-78	91

711	8	700244175H1	SATMON010	g1532072	BLASTN	1044	1e-78	90
712	8	700045296H1	SATMON004	g1532072	BLASTN	1041	1e-77	93
713	8	700578391H1	SATMON031	g1532072	BLASTN	624	1e-76	88
714	8	700346136H1	SATMON021	g1532072	BLASTN	751	1e-76	90
715	8	700457852H1	SATMON029	g1532072	BLASTN	777	1e-76	92
716	8	700570111H1	SATMON030	g1532072	BLASTN	814	1e-75	94
717	8	LIB3066-030-Q1-K1-A12	LIB3066	g1532072	BLASTN	922	1e-75	94
718	8	700149657H1	SATMON007	g1532072	BLASTN	1012	1e-75	92
719	8	LIB3060-043-Q1-K1-B4	LIB3060	g1532072	BLASTN	1013	1e-75	97
720	8	700156752H1	SATMON012	g1532072	BLASTN	1013	1e-75	98
721	8	700454114H1	SATMON029	g1532072	BLASTN	455	1e-74	93
722	8	LIB143-053-Q1-E1-E8	LIB143	g1532072	BLASTN	672	1e-74	98
723	8	700029323H1	SATMON003	g1532072	BLASTN	996	1e-74	99
724	8	700156381H1	SATMON007	g1532072	BLASTN	1002	1e-74	97
725	8	700159603H2	SATMON012	g1532072	BLASTN	1003	1e-74	94
726	8	LIB3060-035-Q1-K1-E5	LIB3060	g1532072	BLASTN	603	1e-72	94
727	8	LIB3069-025-Q1-K1-E12	LIB3069	g1532072	BLASTN	734	1e-72	90
728	8	700166680H1	SATMON013	g1532072	BLASTN	971	1e-72	99
729	8	700171123H1	SATMON013	g1532072	BLASTN	973	1e-72	94
730	8	LIB148-042-Q1-E1-G10	LIB148	g1532072	BLASTN	975	1e-72	100
731	8	700612951H1	SATMON033	g1532072	BLASTN	401	1e-71	96
732	8	700265213H1	SATMON017	g1532072	BLASTN	626	1e-71	92
733	8	LIB189-030-Q1-E1-D11	LIB189	g1532072	BLASTN	740	1e-70	92
734	8	700212877H1	SATMON016	g1532072	BLASTN	918	1e-70	93
735	8	700161545H1	SATMON012	g1532072	BLASTN	621	1e-69	99
736	8	700021259H1	SATMON001	g1532072	BLASTN	940	1e-69	93
737	8	LIB3079-015-Q1-K1-C7	LIB3079	g1532072	BLASTN	508	1e-68	96
738	8	LIB3066-055-Q1-K1-F11	LIB3066	g1532072	BLASTN	832	1e-68	85
739	8	700166771H1	SATMON013	g1532072	BLASTN	922	1e-68	88
740	8	LIB3066-003-Q1-K1-E6	LIB3066	g1532072	BLASTN	927	1e-68	86
741	8	700208131H1	SATMON016	g1532047	BLASTN	747	1e-66	85
742	8	700020688H1	SATMON001	g1532072	BLASTN	905	1e-66	90
743	8	700206405H1	SATMON003	g1532072	BLASTN	905	1e-66	100
744	8	LIB3069-042-Q1-K1-C5	LIB3069	g1532072	BLASTN	891	1e-65	99
745	8	700353835H1	SATMON024	g1532072	BLASTN	896	1e-65	96
746	8	700471533H1	SATMON025	g1532072	BLASTN	496	1e-63	99
747	8	700165425H1	SATMON013	g1532072	BLASTN	850	1e-62	100
748	8	LIB3069-054-Q1-K1-C4	LIB3069	g1532072	BLASTN	613	1e-60	87
749	8	700160896H1	SATMON012	g1532072	BLASTN	746	1e-60	95
750	8	LIB3069-042-Q1-K1-A11	LIB3069	g1532072	BLASTN	840	1e-60	98
751	8	700466625H1	SATMON025	g1532072	BLASTN	635	1e-59	86
752	8	700571770H1	SATMON030	g1532072	BLASTN	820	1e-59	84
753	8	LIB143-024-Q1-E1-D2	LIB143	g1532072	BLASTN	813	1e-58	95
754	8	700453677H1	SATMON028	g1532072	BLASTN	508	1e-57	92
755	8	700193979H1	SATMON014	g1532072	BLASTN	790	1e-57	100
756	8	700467828H1	SATMON025	g1532072	BLASTN	779	1e-56	96
757	8	700150072H1	SATMON007	g1532072	BLASTN	776	1e-55	99
758	8	700802869H1	SATMON036	g1532072	BLASTN	702	1e-54	95
759	8	LIB148-051-Q1-E1-B12	LIB148	g1532072	BLASTN	421	1e-53	92
760	8	700075035H1	SATMON007	g1532072	BLASTN	506	1e-53	93
761	8	700471283H1	SATMON025	g1532072	BLASTN	742	1e-53	91
762	8	700166625H1	SATMON013	g1532072	BLASTN	748	1e-53	99
763	8	700019894H1	SATMON001	g1532072	BLASTN	643	1e-51	89
764	8	700204863H1	SATMON003	g1532072	BLASTN	721	1e-51	99

765	8	700431071H1	SATMONN01	g1532072	BLASTN	366	1e-50	94
766	8	700171582H1	SATMON013	g1532072	BLASTN	492	1e-50	99
767	8	700450579H1	SATMON028	g1532072	BLASTN	338	1e-49	97
768	8	700084149H1	SATMON011	g1532072	BLASTN	672	1e-47	98
769	8	700095070H1	SATMON008	g1532072	BLASTN	662	1e-46	98
770	8	700570827H1	SATMON030	g1532072	BLASTN	344	1e-44	83
771	8	LIB3061-057-Q1-K1-G12	LIB3061	g1532072	BLASTN	378	1e-43	75
772	8	700194918H1	SATMON014	g1532072	BLASTN	626	1e-43	89
773	8	700378894H1	SATMON020	g1532072	BLASTN	495	1e-42	97
774	8	700468029H1	SATMON025	g1532072	BLASTN	616	1e-42	98
775	8	LIB143-012-Q1-E1-H8	LIB143	g1532072	BLASTN	637	1e-42	98
776	8	LIB3060-037-Q1-K1-H4	LIB3060	g1532072	BLASTN	638	1e-42	86
777	8	700433327H1	SATMONN01	g1532072	BLASTN	365	1e-41	86
778	8	700623611H1	SATMON034	g1532072	BLASTN	316	1e-39	95
779	8	700166123H1	SATMON013	g1532072	BLASTN	585	1e-39	100
780	8	700616273H1	SATMON033	g1532072	BLASTN	534	1e-38	98
781	8	700464702H1	SATMON025	g1532072	BLASTN	566	1e-38	99
782	8	700338809H1	SATMON020	g1532072	BLASTN	550	1e-37	100
783	8	700092622H1	SATMON008	g1532072	BLASTN	561	1e-37	99
784	8	LIB3060-043-Q1-K1-A10	LIB3060	g1532072	BLASTN	352	1e-36	93
785	8	700265611H1	SATMON017	g1532072	BLASTN	548	1e-36	95
786	8	700100319H1	SATMON009	g1532072	BLASTN	552	1e-36	98
787	8	700092696H1	SATMON008	g1532072	BLASTN	552	1e-36	98
788	8	700076750H1	SATMON007	g1532072	BLASTN	526	1e-35	99
789	8	700082896H1	SATMON011	g1532072	BLASTN	526	1e-35	99
790	8	700075411H1	SATMON007	g1532072	BLASTN	319	1e-34	91
791	8	701178051H1	SATMONN05	g1532072	BLASTN	342	1e-34	94
792	8	700266358H1	SATMON017	g1532072	BLASTN	520	1e-34	95
793	8	700453981H1	SATMON029	g1532072	BLASTN	505	1e-33	96
794	8	700584238H1	SATMON031	g1532072	BLASTN	509	1e-33	91
795	8	700103871H1	SATMON010	g1532072	BLASTN	491	1e-32	99
796	8	700264460H1	SATMON017	g1532072	BLASTN	495	1e-32	95
797	8	700205122H1	SATMON003	g1532072	BLASTN	501	1e-32	99
798	8	700165768H1	SATMON013	g1532072	BLASTN	420	1e-31	98
799	8	700077179H1	SATMON007	g1532072	BLASTN	471	1e-30	98
800	8	700476654H1	SATMON025	g1532072	BLASTN	476	1e-30	98
801	8	700027374H1	SATMON003	g1532072	BLASTN	476	1e-30	98
802	8	700266994H1	SATMON017	g1532072	BLASTN	476	1e-30	98
803	8	700088193H1	SATMON011	g1532072	BLASTN	488	1e-30	97
804	8	700214511H1	SATMON016	g1532072	BLASTN	298	1e-29	93
805	8	700257186H1	SATMON017	g1532072	BLASTN	465	1e-29	95
806	8	700335814H1	SATMON019	g1532072	BLASTN	469	1e-29	93
807	8	700236166H1	SATMON010	g1532072	BLASTN	450	1e-28	95
808	8	700468243H1	SATMON025	g1532072	BLASTN	456	1e-28	98
809	8	700096327H1	SATMON008	g1532072	BLASTN	451	1e-27	98
810	8	700471139H1	SATMON025	g1532072	BLASTN	451	1e-27	98
811	8	700050420H1	SATMON003	g1532072	BLASTN	365	1e-26	92
812	8	700264846H1	SATMON017	g1532072	BLASTN	430	1e-25	94
813	8	700266803H1	SATMON017	g1532072	BLASTN	293	1e-24	75
814	8	700086134H1	SATMON011	g1532072	BLASTN	422	1e-24	97
815	8	700162001H1	SATMON012	g1532072	BLASTN	279	1e-23	96
816	8	700044825H1	SATMON004	g1532072	BLASTN	411	1e-23	98
817	8	700074679H1	SATMON007	g1532072	BLASTN	411	1e-23	98
818	8	700267694H1	SATMON017	g1532072	BLASTN	320	1e-22	92

819	8	700267990H1	SATMON017	g1532072	BLASTN	361	1e-22	99
820	8	700456715H1	SATMON029	g1532072	BLASTN	293	1e-21	97
821	8	700454806H1	SATMON029	g1532072	BLASTN	356	1e-21	98
822	8	700466812H1	SATMON025	g1532072	BLASTN	380	1e-21	95
823	8	700046476H1	SATMON004	g1532072	BLASTN	381	1e-21	97
824	8	700207160H1	SATMON017	g1532072	BLASTN	393	1e-21	96
825	8	700383037H1	SATMON024	g1532072	BLASTN	280	1e-20	98
826	8	700549660H1	SATMON022	g1532072	BLASTN	376	1e-20	97
827	8	LIB3066-004-Q1-K1-G12	LIB3066	g1532072	BLASTN	379	1e-20	91
828	8	700801422H1	SATMON036	g1532072	BLASTN	321	1e-19	97
829	8	700045319H1	SATMON004	g1532072	BLASTN	336	1e-17	99
830	8	700046047H1	SATMON004	g1532072	BLASTN	336	1e-17	99
831	8	700264328H1	SATMON017	g1532072	BLASTN	339	1e-17	95
832	8	700083485H1	SATMON011	g1532072	BLASTN	341	1e-17	99
833	8	700461268H1	SATMON033	g1532072	BLASTN	342	1e-17	95
834	8	700053271H1	SATMON008	g1532072	BLASTN	311	1e-15	98
835	8	700215275H1	SATMON016	g1532072	BLASTN	321	1e-15	98
836	8	700623133H1	SATMON034	g1532072	BLASTN	275	1e-14	75
837	8	700454949H1	SATMON029	g1532072	BLASTN	280	1e-14	99
838	8	700440763H1	SATMON026	g1532072	BLASTN	307	1e-14	95
839	8	LIB3068-022-Q1-K1-C4	LIB3068	g1532072	BLASTN	150	1e-12	94
840	8	700798849H1	SATMON036	g1532072	BLASTN	281	1e-12	98
841	8	700333057H1	SATMON019	g1532072	BLASTN	184	1e-11	91
842	8	700265786H1	SATMON017	g1532072	BLASTN	200	1e-11	90
843	8	700193030H1	SATMON014	g1532072	BLASTN	158	1e-9	94
844	8	700262964H1	SATMON017	g1532047	BLASTN	171	1e-9	81
845	8	LIB3069-023-Q1-K1-B2	LIB3069	g1403043	BLASTN	195	1e-9	84
846	8	700474096H1	SATMON025	g1532072	BLASTN	199	1e-9	97
847	8	700028326H1	SATMON003	g1532072	BLASTN	151	1e-8	96
848	8011	700440327H1	SATMON026	g1532047	BLASTN	292	1e-14	85
849	851	LIB3059-052-Q1-K1-E6	LIB3059	g1403043	BLASTN	1033	1e-85	75
850	851	700090958H1	SATMON011	g1532072	BLASTN	1044	1e-78	80
851	851	700224605H1	SATMON011	g1532072	BLASTN	850	1e-62	80
852	851	700551562H1	SATMON022	g1403043	BLASTN	374	1e-55	79
853	851	700153939H1	SATMON007	g1403043	BLASTN	772	1e-55	80
854	851	700469231H1	SATMON025	g1532072	BLASTN	560	1e-52	76
855	851	700349854H1	SATMON023	g1403043	BLASTN	669	1e-46	81
856	851	701169324H1	SATMONN05	g1403043	BLASTN	669	1e-46	78
857	851	700088319H1	SATMON011	g1532072	BLASTN	492	1e-30	78
2480	-700998660	700998660H1	SOYMON018	g1531764	BLASTN	249	1e-29	90
2481	-GM927	LIB3028-005-Q1-B1-B4	LIB3028	g1421750	BLASTN	261	1e-10	80
2482	13379	700842514H1	SOYMON020	g1915980	BLASTN	575	1e-41	73
2483	13379	701042786H1	SOYMON029	g1421750	BLASTN	377	1e-20	80
2484	15556	701101584H1	SOYMON028	g1421750	BLASTN	526	1e-35	69
2485	15556	701099749H1	SOYMON028	g1155239	BLASTN	453	1e-27	72
2486	16	LIB3051-018-Q1-E1-A7	LIB3051	g1421750	BLASTN	1146	1e-93	81
2487	16	LIB3056-005-Q1-N1-E10	LIB3056	g1421750	BLASTN	944	1e-88	80
2488	16	LIB3051-101-Q1-K1-C7	LIB3051	g1421750	BLASTN	1135	1e-85	80
2489	16	LIB3056-001-Q1-B1-H12	LIB3056	g1421750	BLASTN	752	1e-82	81
2490	16	700661334H1	SOYMON005	g1421750	BLASTN	1092	1e-82	80
2491	16	LIB3056-012-Q1-N1-B12	LIB3056	g1421750	BLASTN	1085	1e-81	80
2492	16	LIB3055-011-Q1-N1-G2	LIB3055	g1421750	BLASTN	958	1e-78	79
2493	16	700663981H1	SOYMON005	g1421750	BLASTN	996	1e-74	82
2494	16	701109737H1	SOYMON036	g1421750	BLASTN	955	1e-70	83

2495	16	701130190H1	SOYMON037	g1421750	BLASTN	935	1e-69	81
2496	16	701003301H1	SOYMON019	g1421750	BLASTN	412	1e-68	82
2497	16	700894137H1	SOYMON024	g1421750	BLASTN	923	1e-68	84
2498	16	700980096H1	SOYMON009	g1421750	BLASTN	923	1e-68	81
2499	16	700942625H1	SOYMON024	g1421750	BLASTN	911	1e-67	80
2500	16	700829695H1	SOYMON019	g1421750	BLASTN	918	1e-67	84
2501	16	700662572H1	SOYMON005	g1421750	BLASTN	920	1e-67	84
2502	16	701127647H1	SOYMON037	g1421750	BLASTN	902	1e-66	84
2503	16	701014550H1	SOYMON019	g1421750	BLASTN	556	1e-65	81
2504	16	LIB3051-043-Q1-K1-G3	LIB3051	g1421750	BLASTN	698	1e-65	82
2505	16	700730914H1	SOYMON009	g1421750	BLASTN	897	1e-65	84
2506	16	701052455H1	SOYMON032	g1421750	BLASTN	876	1e-64	81
2507	16	LIB3051-074-Q1-K1-A7	LIB3051	g1421750	BLASTN	876	1e-64	75
2508	16	700874839H1	SOYMON018	g1421750	BLASTN	862	1e-63	82
2509	16	701005943H1	SOYMON019	g1421750	BLASTN	865	1e-63	82
2510	16	700663357H1	SOYMON005	g1421750	BLASTN	869	1e-63	82
2511	16	701042053H1	SOYMON029	g1421750	BLASTN	872	1e-63	81
2512	16	700971988H1	SOYMON005	g1421750	BLASTN	508	1e-61	82
2513	16	701010728H1	SOYMON019	g1421750	BLASTN	657	1e-61	81
2514	16	700987206H1	SOYMON009	g1421750	BLASTN	845	1e-61	83
2515	16	701126773H1	SOYMON037	g1421750	BLASTN	847	1e-61	79
2516	16	700764714H1	SOYMON023	g1421750	BLASTN	830	1e-60	81
2517	16	700974444H1	SOYMON005	g1421750	BLASTN	830	1e-60	81
2518	16	701118620H1	SOYMON037	g1421750	BLASTN	832	1e-60	75
2519	16	700729926H1	SOYMON009	g1421750	BLASTN	834	1e-60	79
2520	16	700945142H1	SOYMON024	g1421750	BLASTN	470	1e-59	84
2521	16	700867912H1	SOYMON016	g1421750	BLASTN	820	1e-59	82
2522	16	700746546H1	SOYMON013	g1421750	BLASTN	824	1e-59	78
2523	16	700873331H1	SOYMON018	g1421750	BLASTN	769	1e-58	82
2524	16	700738212H1	SOYMON012	g1421750	BLASTN	780	1e-56	83
2525	16	700830454H1	SOYMON019	g1421750	BLASTN	789	1e-56	81
2526	16	700726129H1	SOYMON009	g1421750	BLASTN	789	1e-56	81
2527	16	700996774H1	SOYMON018	g1421750	BLASTN	486	1e-55	81
2528	16	700746427H1	SOYMON013	g1421750	BLASTN	766	1e-55	77
2529	16	701123911H1	SOYMON037	g1421750	BLASTN	769	1e-55	76
2530	16	700745730H1	SOYMON013	g1421750	BLASTN	775	1e-55	76
2531	16	700900940H1	SOYMON027	g1421750	BLASTN	760	1e-54	78
2532	16	700846419H1	SOYMON021	g1421750	BLASTN	762	1e-54	78
2533	16	701203427H1	SOYMON035	g1421750	BLASTN	491	1e-53	84
2534	16	LIB3051-074-Q1-K1-F5	LIB3051	g1421750	BLASTN	650	1e-53	75
2535	16	700747646H1	SOYMON013	g1421750	BLASTN	747	1e-53	77
2536	16	701049823H1	SOYMON032	g1421750	BLASTN	751	1e-53	77
2537	16	701049188H1	SOYMON032	g1421750	BLASTN	731	1e-52	75
2538	16	700875156H1	SOYMON018	g1421750	BLASTN	737	1e-52	78
2539	16	700864540H1	SOYMON016	g1421750	BLASTN	741	1e-52	81
2540	16	700682329H2	SOYMON008	g1421750	BLASTN	719	1e-51	76
2541	16	700846215H1	SOYMON021	g1421750	BLASTN	575	1e-50	82
2542	16	701101327H1	SOYMON028	g1421750	BLASTN	592	1e-50	77
2543	16	LIB3055-011-Q1-N1-E7	LIB3055	g1421750	BLASTN	680	1e-50	80
2544	16	700848992H1	SOYMON021	g1421750	BLASTN	708	1e-50	76
2545	16	700966820H1	SOYMON028	g1421750	BLASTN	710	1e-50	75
2546	16	701051785H1	SOYMON032	g1421750	BLASTN	717	1e-50	75
2547	16	700681079H1	SOYMON008	g1917012	BLASTN	290	1e-49	77
2548	16	701138880H1	SOYMON038	g1421750	BLASTN	610	1e-49	78

2549	16	700872967H1	SOYMON018	g1421750	BLASTN	704	1e-49	75
2550	16	701119960H1	SOYMON037	g1421750	BLASTN	684	1e-48	81
2551	16	700873306H1	SOYMON018	g1421750	BLASTN	693	1e-48	83
2552	16	700751484H1	SOYMON014	g1421750	BLASTN	635	1e-47	82
2553	16	700958865H1	SOYMON022	g1421750	BLASTN	486	1e-46	82
2554	16	700872833H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2555	16	700871673H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2556	16	700872801H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2557	16	700847572H1	SOYMON021	g1421750	BLASTN	660	1e-46	75
2558	16	700728046H1	SOYMON009	g1421750	BLASTN	661	1e-46	84
2559	16	700894721H1	SOYMON024	g1421750	BLASTN	550	1e-43	84
2560	16	700730946H1	SOYMON009	g1421750	BLASTN	307	1e-42	77
2561	16	700727943H1	SOYMON009	g1421750	BLASTN	378	1e-42	79
2562	16	701101238H1	SOYMON028	g1421750	BLASTN	456	1e-42	92
2563	16	700758519H1	SOYMON015	g1421750	BLASTN	358	1e-41	77
2564	16	700740343H1	SOYMON012	g1531764	BLASTN	607	1e-41	72
2565	16	700848076H1	SOYMON021	g1531764	BLASTN	469	1e-40	89
2566	16	LIB3051-080-Q1-K1-H7	LIB3051	g1531764	BLASTN	472	1e-38	90
2567	16	700955916H1	SOYMON022	g1531764	BLASTN	427	1e-37	90
2568	16	700873876H1	SOYMON018	g1421750	BLASTN	465	1e-37	80
2569	16	700746271H1	SOYMON013	g1421750	BLASTN	515	1e-37	80
2570	16	700662887H1	SOYMON005	g1421750	BLASTN	506	1e-35	82
2571	16	700983591H1	SOYMON009	g1421750	BLASTN	537	1e-35	82
2572	16	700752343H1	SOYMON014	g1421750	BLASTN	488	1e-34	73
2573	16	700901973H1	SOYMON027	g1421750	BLASTN	490	1e-34	82
2574	16	LIB3053-013-Q1-N1-B11	LIB3053	g1421750	BLASTN	522	1e-34	75
2575	16	700988481H1	SOYMON009	g1421750	BLASTN	506	1e-33	78
2576	16	LIB3051-078-Q1-K1-C12	LIB3051	g1421750	BLASTN	302	1e-32	75
2577	16	700985640H1	SOYMON009	g1421750	BLASTN	449	1e-30	86
2578	16	701127038H1	SOYMON037	g1421750	BLASTN	467	1e-30	67
2579	16	700898677H1	SOYMON027	g1531764	BLASTN	474	1e-30	89
2580	16	700742260H1	SOYMON012	g1421750	BLASTN	476	1e-30	70
2581	16	701103203H1	SOYMON028	g1421750	BLASTN	287	1e-27	81
2582	16	700897144H1	SOYMON027	g1421750	BLASTN	264	1e-26	75
2583	16	700832039H1	SOYMON019	g1421750	BLASTN	390	1e-26	82
2584	16	700743311H1	SOYMON012	g1421750	BLASTN	429	1e-26	87
2585	16	LIB3040-044-Q1-E1-B8	LIB3040	g1421750	BLASTN	282	1e-25	92
2586	16	LIB3051-024-Q1-K1-C4	LIB3051	g2394382	BLASTX	110	1e-24	95
2587	16	701011765H1	SOYMON019	g1421750	BLASTN	383	1e-21	76
2588	16	700893428H1	SOYMON024	g1421752	BLASTX	106	1e-20	82
2589	16	700996204H1	SOYMON018	g1421750	BLASTN	288	1e-20	75
2590	16	LIB3049-028-Q1-E1-C6	LIB3049	g1421750	BLASTN	230	1e-19	90
2591	16	701103628H1	SOYMON028	g1421750	BLASTN	277	1e-18	73
2592	16	700972636H1	SOYMON005	g1421750	BLASTN	286	1e-16	73
2593	16	700875417H1	SOYMON018	g1421750	BLASTN	209	1e-15	75
2594	16	701207702H1	SOYMON035	g1421752	BLASTX	153	1e-14	83
2595	16	701054556H1	SOYMON032	g1421750	BLASTN	230	1e-14	75
2596	16	701046407H1	SOYMON032	g1421750	BLASTN	230	1e-14	75
2597	16	701117634H1	SOYMON037	g1421750	BLASTN	230	1e-14	74
2598	16	701127714H1	SOYMON037	g1421750	BLASTN	230	1e-14	75
2599	16	700663239H1	SOYMON005	g1421750	BLASTN	210	1e-13	74
2600	16	700738510H1	SOYMON012	g1421750	BLASTN	212	1e-13	72
2601	16	700943559H1	SOYMON024	g1421750	BLASTN	215	1e-13	71
2602	16	700666278H1	SOYMON005	g1421750	BLASTN	290	1e-13	93

2603	16	700663144H1	SOYMON005	g1421750	BLASTN	204	1e-12	74
2604	16	700684147H1	SOYMON008	g1421750	BLASTN	278	1e-12	89
2605	16	700953076H1	SOYMON022	g1421750	BLASTN	285	1e-12	92
2606	16	701099931H1	SOYMON028	g1421750	BLASTN	202	1e-11	72
2607	16	701058910H1	SOYMON033	g1421750	BLASTN	216	1e-11	74
2608	16	700754949H1	SOYMON014	g1421750	BLASTN	266	1e-11	89
2609	16	700562967H1	SOYMON002	g1421750	BLASTN	266	1e-11	82
2610	16	701012790H1	SOYMON019	g1421750	BLASTN	251	1e-10	89
2611	16	701101739H1	SOYMON028	g1421750	BLASTN	260	1e-10	82
2612	16	701118211H1	SOYMON037	g1490554	BLASTX	97	1e-9	47
2613	16	700844394H1	SOYMON021	g1917013	BLASTX	116	1e-9	85
2614	16	701048560H1	SOYMON032	g1421750	BLASTN	198	1e-9	79
2615	16	701062373H1	SOYMON033	g1421750	BLASTN	203	1e-9	71
2616	16	701120496H1	SOYMON037	g1421750	BLASTN	230	1e-9	82
2617	16	701049680H1	SOYMON032	g1421750	BLASTN	177	1e-8	70
2618	16	700748760H1	SOYMON013	g1421750	BLASTN	230	1e-8	91
2619	16	700889482H1	SOYMON024	g1421750	BLASTN	235	1e-8	91
2620	16	700725013H1	SOYMON009	g1421750	BLASTN	239	1e-8	85
2621	16048	LIB3028-004-Q1-B1-E2	LIB3028	g1421750	BLASTN	879	1e-64	68
2622	16048	700761667H1	SOYMON015	g1421750	BLASTN	528	1e-35	71
2623	16048	700958003H1	SOYMON022	g1421750	BLASTN	428	1e-26	78

ASPARTATE KINASE (EC 2.7.2.4)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
858	-700018870	700018870H1	SATMON001	g500850	BLASTN	1095	1e-82	100
859	-700085903	700085903H1	SATMON011	g500852	BLASTN	1606	1e-124	99
860	-700086169	700086169H1	SATMON011	g500852	BLASTN	559	1e-94	92
861	-700096679	700096679H1	SATMON008	g500850	BLASTN	1131	1e-85	99
862	-700096794	700096794H1	SATMON008	g500850	BLASTN	1515	1e-117	100
863	-700106390	700106390H1	SATMON010	g2243115	BLASTN	544	1e-51	72
864	-700168286	700168286H1	SATMON013	g2243115	BLASTN	519	1e-34	72
865	-700171363	700171363H1	SATMON013	g500850	BLASTN	1085	1e-81	94
866	-700194781	700194781H1	SATMON014	g2257742	BLASTN	635	1e-44	79
867	-700213839	700213839H1	SATMON016	g500850	BLASTN	664	1e-75	94
868	-700219756	700219756H1	SATMON011	g500850	BLASTN	1359	1e-104	99
869	-700258808	700258808H1	SATMON017	g500850	BLASTN	630	1e-85	96
870	-700263439	700263439H1	SATMON017	g2243115	BLASTN	448	1e-43	76
871	-700266615	700266615H1	SATMON017	g2243116	BLASTX	179	1e-17	73
872	-700342655	700342655H1	SATMON021	g2257743	BLASTX	213	1e-22	82
873	-700343285	700343285H1	SATMON021	g2257742	BLASTN	437	1e-25	66
874	-700467533	700467533H1	SATMON025	g2243115	BLASTN	439	1e-26	79
875	-700548678	700548678H1	SATMON022	g147979	BLASTX	147	1e-13	62
876	-700613618	700613618H1	SATMON033	g500850	BLASTN	965	1e-81	98
877	-L30691987	LIB3069-018-Q1-K1-A3	LIB3069	g2243115	BLASTN	266	1e-10	57
878	12201	700457103H1	SATMON029	g2257742	BLASTN	789	1e-56	76
879	12201	700457111H1	SATMON029	g2243115	BLASTN	513	1e-33	76
880	12931	700380864H1	SATMON023	g500852	BLASTN	1450	1e-111	95
881	12931	700105610H1	SATMON010	g500852	BLASTN	1046	1e-105	95
882	12931	700380848H1	SATMON023	g500852	BLASTN	1193	1e-96	95
883	12931	700205392H1	SATMON003	g500852	BLASTN	1065	1e-94	93
884	12931	700552314H1	SATMON022	g500852	BLASTN	908	1e-86	90
885	12931	700551915H1	SATMON022	g500852	BLASTN	1090	1e-81	90

886	16037	700344509H1	SATMON021	g500852	BLASTN	1184	1e-89	92
887	16037	700345170H1	SATMON021	g500852	BLASTN	600	1e-58	85
888	16157	700212607H1	SATMON016	g2243115	BLASTN	914	1e-67	76
889	16157	700094809H1	SATMON008	g2243116	BLASTX	144	1e-12	84
890	19231	700091761H1	SATMON011	g500850	BLASTN	1358	1e-104	98
891	19231	700612568H1	SATMON033	g500850	BLASTN	1137	1e-102	99
892	22303	700553291H1	SATMON022	g2243115	BLASTN	644	1e-44	70
893	22303	700553382H1	SATMON022	g2243115	BLASTN	418	1e-39	70
894	28000	LIB143-061-Q1-E1-C7	LIB143	g2243115	BLASTN	1132	1e-85	75
895	28000	700474110H1	SATMON025	g2243115	BLASTN	509	1e-33	75
896	30401	700620948H1	SATMON034	g500852	BLASTN	327	1e-30	88
897	32907	LIB143-038-Q1-E1-B11	LIB143	g500850	BLASTN	1864	1e-146	96
898	32907	700096779H1	SATMON008	g500850	BLASTN	1490	1e-115	97
899	5616	700346488H1	SATMON021	g2243115	BLASTN	664	1e-46	71
900	5616	700196138H1	SATMON014	g2243115	BLASTN	630	1e-43	74
2624	-700556108	700556108H1	SOYMON001	g2243115	BLASTN	700	1e-49	74
2625	-700663367	700663367H1	SOYMON005	g2243115	BLASTN	737	1e-52	77
2626	-700733301	700733301H1	SOYMON010	g2243115	BLASTN	751	1e-53	78
2627	-700747979	700747979H1	SOYMON013	g2257742	BLASTN	449	1e-27	70
2628	-700832664	700832664H1	SOYMON019	g167547	BLASTN	322	1e-44	78
2629	-700843925	700843925H1	SOYMON021	g167547	BLASTN	616	1e-42	71
2630	-700888516	700888516H1	SOYMON024	g464225	BLASTX	193	1e-19	78
2631	-700892002	700892002H1	SOYMON024	g2243115	BLASTN	363	1e-21	79
2632	-700959057	700959057H1	SOYMON022	g2257742	BLASTN	497	1e-32	72
2633	-700971891	700971891H1	SOYMON005	g167547	BLASTN	699	1e-49	74
2634	-700984812	700984812H1	SOYMON009	g2257742	BLASTN	801	1e-57	77
2635	-701069254	701069254H1	SOYMON034	g2243115	BLASTN	260	1e-23	75
2636	-701120341	701120341H1	SOYMON037	g2243115	BLASTN	567	1e-38	77
2637	-GM35173	LIB3051-037-Q1-K1-B9	LIB3051	g2970554	BLASTN	193	1e-11	83
2638	15020	700557507H1	SOYMON001	g167547	BLASTN	914	1e-67	79
2639	15020	700666142H1	SOYMON005	g1107460	BLASTN	767	1e-55	77
2640	18237	700797368H1	SOYMON017	g2257742	BLASTN	819	1e-59	81
2641	18237	700797360H1	SOYMON017	g2257742	BLASTN	809	1e-58	84
2642	19332	LIB3056-004-Q1-N1-D5	LIB3056	g2243115	BLASTN	1118	1e-84	74
2643	19332	700786255H2	SOYMON011	g2257742	BLASTN	626	1e-43	72
2644	19332	700684751H1	SOYMON008	g2257742	BLASTN	583	1e-39	73
2645	21954	701100440H1	SOYMON028	g167547	BLASTN	835	1e-60	78
2646	21954	701059173H1	SOYMON033	g167547	BLASTN	719	1e-51	75
2647	26336	701003103H1	SOYMON019	g2243115	BLASTN	877	1e-64	79
2648	26336	700976874H1	SOYMON009	g2243115	BLASTN	880	1e-64	79

ASPARTATE-SEMIALDEHYDE DEHYDROGENASE (EC 1.2.1.11)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
901	-700439614	700439614H1	SATMON026	g2314350	BLASTX	107	1e-13	52
902	-701183695	701183695H1	SATMONN06	g289910	BLASTX	150	1e-13	69
903	-L1487398	LIB148-064-Q1-E1-D10	LIB148	g1749466	BLASTX	178	1e-33	55
904	-L30622830	LIB3062-028-Q1-K1-A8	LIB3062	g1085109	BLASTX	108	1e-40	48
2649	-700756763	700756763H1	SOYMON014	g1359593	BLASTX	71	1e-8	52
2650	-700830054	700830054H1	SOYMON019	g142828	BLASTX	87	1e-10	41
2651	-701105617	701105617H1	SOYMON036	g142828	BLASTX	215	1e-22	57
2652	-GM8539	LIB3039-047-Q1-E1-C6	LIB3039	g142828	BLASTX	188	1e-35	50
2653	30187	LIB3049-001-Q1-E1-F2	LIB3049	g1359593	BLASTX	136	1e-26	54

2654 30187 700556105H1 SOYMON001 g1359593 BLASTX 132 1e-11 56

O-SUCCINYLMOMOSERINE (THIOL)-LYASE (EC 4.2.99.9)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
905	-700049526	700049526H1	SATMON003	g2198852	BLASTN	251	1e-9	76
906	-700086788	700086788H1	SATMON011	g2198850	BLASTN	631	1e-61	88
907	-700460589	700460589H1	SATMON030	g2198850	BLASTN	251	1e-32	83
908	-700577561	700577561H1	SATMON031	g2198852	BLASTN	207	1e-15	82
909	-700579840	700579840H1	SATMON031	g2198850	BLASTN	216	1e-12	94
910	-700616013	700616013H1	SATMON033	g2198852	BLASTN	278	1e-39	81
911	-L1892785	LIB189-011-Q1-E1-A10	LIB189	g2198852	BLASTN	311	1e-14	82
912	-L30594402	LIB3059-042-Q1-K1-E11	LIB3059	g2198852	BLASTN	489	1e-40	84
913	-L30604293	LIB3060-028-Q1-K1-E7	LIB3060	g2198852	BLASTN	380	1e-20	83
914	-L30693289	LIB3069-016-Q1-K1-G10	LIB3069	g2198852	BLASTN	197	1e-10	79
915	10571	700224881H1	SATMON011	g2198852	BLASTN	319	1e-15	75
916	10801	700429049H1	SATMONN01	g2198852	BLASTN	210	1e-16	82
917	10801	700167813H1	SATMON013	g2198852	BLASTN	200	1e-15	82
918	10801	700074027H1	SATMON007	g2198852	BLASTN	178	1e-11	80
919	16379	LIB3061-035-Q1-K1-B6	LIB3061	g2198850	BLASTN	2184	1e-173	99
920	16379	700259309H1	SATMON017	g2198850	BLASTN	1259	1e-108	92
921	16379	700051696H1	SATMON003	g2198850	BLASTN	1392	1e-107	96
922	16379	700239553H1	SATMON010	g2198850	BLASTN	1265	1e-96	96
923	16379	700042846H1	SATMON004	g2198850	BLASTN	1208	1e-91	95
924	16379	700092741H1	SATMON008	g2198850	BLASTN	1042	1e-89	95
925	16379	LIB3061-017-Q1-K1-D7	LIB3061	g2198850	BLASTN	1183	1e-89	99
926	16379	700206765H1	SATMON003	g2198850	BLASTN	1095	1e-82	81
927	16379	700150281H1	SATMON007	g2198850	BLASTN	1024	1e-76	94
928	16379	700165862H1	SATMON013	g2198850	BLASTN	948	1e-75	94
929	2221	LIB3060-017-Q1-K1-B10	LIB3060	g2198850	BLASTN	1819	1e-161	99
930	2221	LIB84-006-Q1-E1-F3	LIB84	g2198852	BLASTN	1615	1e-153	97
931	2221	700575334H1	SATMON030	g2198850	BLASTN	1570	1e-124	99
932	2221	700206250H1	SATMON003	g2198850	BLASTN	1515	1e-117	100
933	2221	700095073H1	SATMON008	g2198852	BLASTN	1490	1e-115	100
934	2221	700571230H1	SATMON030	g2198850	BLASTN	1355	1e-114	93
935	2221	700157358H1	SATMON012	g2198850	BLASTN	1370	1e-105	100
936	2221	700379811H1	SATMON021	g2198850	BLASTN	1345	1e-103	93
937	2221	700041570H1	SATMON004	g2198850	BLASTN	1325	1e-101	100
938	2221	700104063H1	SATMON010	g2198850	BLASTN	1157	1e-95	90
939	2221	700378265H1	SATMON019	g2198850	BLASTN	771	1e-94	99
940	2221	700235329H1	SATMON010	g2198850	BLASTN	1234	1e-94	93
941	2221	LIB3068-057-Q1-K1-D1	LIB3068	g2198852	BLASTN	1209	1e-91	94
942	2221	700159158H1	SATMON012	g2198850	BLASTN	1174	1e-89	94
943	2221	700580854H1	SATMON031	g2198850	BLASTN	754	1e-86	92
944	2221	700623409H1	SATMON034	g2198850	BLASTN	910	1e-84	96
945	2221	701164706H1	SATMONN04	g2198852	BLASTN	577	1e-83	94
946	2221	700164719H1	SATMON013	g2198852	BLASTN	831	1e-83	99
947	2221	700158146H1	SATMON012	g2198850	BLASTN	1100	1e-82	93
948	2221	700203970H1	SATMON003	g2198852	BLASTN	996	1e-80	99
949	2221	700158313H1	SATMON012	g2198850	BLASTN	1036	1e-77	93
950	2221	700425211H1	SATMONN01	g2198852	BLASTN	485	1e-61	96
951	2221	700167764H1	SATMON013	g2198850	BLASTN	793	1e-57	93
952	23788	700102780H1	SATMON010	g2198852	BLASTN	1131	1e-104	99

953	23788	701167458H1	SATMONN05	g2198852	BLASTN	1069	1e-90	93
2655	-700900206	700900206H1	SOYMON027	g1742961	BLASTX	215	1e-24	78
2656	-GM40351	LIB3051-114-Q1-K1-H12	LIB3051	g2198851	BLASTX	122	1e-25	96
2657	12502	701101592H1	SOYMON028	g146846	BLASTX	103	1e-18	44
2658	12502	701106834H1	SOYMON036	g146846	BLASTX	103	1e-18	44
2659	13820	LIB3055-003-Q1-N1-F12	LIB3055	g3202028	BLASTX	193	1e-35	94
2660	8119	700989656H1	SOYMON011	g1742960	BLASTN	815	1e-59	79

CYSTATHIONINE β -LYASE (EC 4.4.1.8)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
954	-700155172	700155172H1	SATMON007	g704397	BLASTX	361	1e-43	78
955	-L362943	LIB36-013-Q1-E1-G10	LIB36	g704396	BLASTN	818	1e-59	75
956	19856	700240664H1	SATMON010	g704396	BLASTN	496	1e-31	69
957	19856	700572496H1	SATMON030	g704396	BLASTN	446	1e-26	68
958	22960	701170964H1	SATMONN05	g704396	BLASTN	778	1e-56	78
959	22960	LIB3061-002-Q1-K2-F9	LIB3061	g704396	BLASTN	765	1e-53	77
960	22960	701172780H2	SATMONN05	g704396	BLASTN	686	1e-48	73
961	22960	700578571H1	SATMON031	g704397	BLASTX	222	1e-23	89
962	30752	LIB3078-055-Q1-K1-C8	LIB3078	g704396	BLASTN	747	1e-51	71
963	30752	700086603H1	SATMON011	g704397	BLASTX	192	1e-18	64
2661	-701001147	701001147H1	SOYMON018	g704396	BLASTN	847	1e-61	78
2662	18602	700566066H1	SOYMON002	g704396	BLASTN	751	1e-57	79
2663	18602	700890955H1	SOYMON024	g704396	BLASTN	698	1e-49	77
2664	18602	700896865H1	SOYMON027	g704396	BLASTN	682	1e-48	77
2665	5144	LIB3050-006-Q1-E1-A9	LIB3050	g1399263	BLASTX	96	1e-31	41

5-METHYLTETRAHYDROPTEROYLTRIGLUTAMATE--HOMOCYSTEINE S-METHYLTRANSFERASE (EC 2.1.1.14)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
964	-700212217	700212217H1	SATMON016	g886471	BLASTX	151	1e-16	93
965	-700333966	700333966H1	SATMON019	g886470	BLASTN	370	1e-23	66
966	-700377403	700377403H1	SATMON019	g2738247	BLASTN	305	1e-26	80
967	-700571893	700571893H1	SATMON030	g974781	BLASTN	262	1e-20	80
968	-701165656	701165656H1	SATMONN04	g2738247	BLASTN	450	1e-50	73
969	-L30622954	LIB3062-030-Q1-K1-C9	LIB3062	g886470	BLASTN	366	1e-26	81
970	-L30662004	LIB3066-026-Q1-K1-F1	LIB3066	g2738248	BLASTX	140	1e-28	77
971	-L30671450	LIB3067-001-Q1-K1-C6	LIB3067	g2738247	BLASTN	553	1e-46	69
972	-L30694410	LIB3069-057-Q1-K1-B4	LIB3069	g886470	BLASTN	447	1e-26	67
973	1	700452404H1	SATMON028	g453939	BLASTX	59	1e-17	94
974	13513	700049012H1	SATMON003	g1814402	BLASTN	1006	1e-74	81
975	13513	700086058H1	SATMON011	g1814402	BLASTN	982	1e-72	81
976	13513	700235814H1	SATMON010	g1814402	BLASTN	890	1e-65	79
977	13513	700170015H1	SATMON013	g1814402	BLASTN	578	1e-39	78
978	3835	700093161H1	SATMON008	g974781	BLASTN	858	1e-62	76
979	3835	700223321H1	SATMON011	g974781	BLASTN	836	1e-60	80
980	3835	700454003H1	SATMON029	g974781	BLASTN	823	1e-59	82
981	3835	700238925H1	SATMON010	g2738247	BLASTN	733	1e-55	74
982	3835	700075142H1	SATMON007	g2738247	BLASTN	569	1e-52	73
983	3835	700151969H1	SATMON007	g974781	BLASTN	660	1e-46	76
984	3835	700281434H2	SATMON019	g974781	BLASTN	560	1e-45	79

985	3835	700084914H1	SATMON011	g974781	BLASTN	440	1e-36	79
986	3835	700215135H1	SATMON016	g974781	BLASTN	543	1e-36	78
987	3835	700202218H1	SATMON003	g974781	BLASTN	389	1e-23	78
988	3835	700281467H2	SATMON019	g2738248	BLASTX	123	1e-12	71
989	456	LIB3059-019-Q1-K1-G6	LIB3059	g974781	BLASTN	1493	1e-115	83
990	456	LIB3068-012-Q1-K1-G7	LIB3068	g974781	BLASTN	1471	1e-113	81
991	456	LIB3061-050-Q1-K1-G10	LIB3061	g886470	BLASTN	1285	1e-98	82
992	456	LIB3067-043-Q1-K1-F9	LIB3067	g2738247	BLASTN	1287	1e-98	79
993	456	700087169H1	SATMON011	g1814402	BLASTN	1245	1e-94	86
994	456	LIB3069-030-Q1-K1-G9	LIB3069	g1814402	BLASTN	772	1e-93	80
995	456	LIB3069-019-Q1-K1-G11	LIB3069	g1814402	BLASTN	1202	1e-91	81
996	456	700202514H1	SATMON003	g1814402	BLASTN	1134	1e-89	84
997	456	LIB3062-011-Q1-K1-F2	LIB3062	g886470	BLASTN	664	1e-86	84
998	456	LIB143-015-Q1-E1-A1	LIB143	g974781	BLASTN	1143	1e-86	81
999	456	700570326H1	SATMON030	g974781	BLASTN	1089	1e-85	83
1000	456	700103727H1	SATMON010	g2738247	BLASTN	1132	1e-85	86
1001	456	700574823H1	SATMON030	g1814402	BLASTN	1135	1e-85	80
1002	456	700091666H1	SATMON011	g1814402	BLASTN	1136	1e-85	84
1003	456	LIB3069-033-Q1-K1-E9	LIB3069	g2738247	BLASTN	1115	1e-84	79
1004	456	700572634H1	SATMON030	g1814402	BLASTN	1123	1e-84	85
1005	456	700211829H1	SATMON016	g2738247	BLASTN	1096	1e-82	84
1006	456	700087109H1	SATMON011	g1814402	BLASTN	1102	1e-82	85
1007	456	LIB3059-003-Q1-K1-B1	LIB3059	g1814402	BLASTN	961	1e-81	82
1008	456	700047556H1	SATMON003	g2738247	BLASTN	976	1e-81	82
1009	456	700201925H1	SATMON003	g2738247	BLASTN	1001	1e-81	84
1010	456	700209343H1	SATMON016	g1814402	BLASTN	1081	1e-81	82
1011	456	700348416H1	SATMON023	g1814402	BLASTN	1081	1e-81	83
1012	456	700095559H1	SATMON008	g1814402	BLASTN	1085	1e-81	82
1013	456	700073472H1	SATMON007	g1814402	BLASTN	1089	1e-81	82
1014	456	LIB3062-057-Q1-K1-C2	LIB3062	g886470	BLASTN	1073	1e-80	78
1015	456	700093445H1	SATMON008	g1814402	BLASTN	1076	1e-80	84
1016	456	LIB143-026-Q1-E1-H3	LIB143	g2738247	BLASTN	648	1e-79	79
1017	456	700206240H1	SATMON003	g1814402	BLASTN	1057	1e-79	82
1018	456	700091907H1	SATMON011	g886470	BLASTN	1062	1e-79	80
1019	456	700258178H1	SATMON017	g1814402	BLASTN	1064	1e-79	84
1020	456	700086565H1	SATMON011	g1814402	BLASTN	1065	1e-79	82
1021	456	700471452H1	SATMON025	g2738247	BLASTN	1045	1e-78	85
1022	456	700243858H1	SATMON010	g1814402	BLASTN	1048	1e-78	85
1023	456	700352224H1	SATMON023	g2738247	BLASTN	1051	1e-78	84
1024	456	700084769H1	SATMON011	g1814402	BLASTN	1054	1e-78	83
1025	456	700082818H1	SATMON011	g886470	BLASTN	984	1e-77	84
1026	456	700331974H1	SATMON019	g886470	BLASTN	1031	1e-77	82
1027	456	700086203H1	SATMON011	g1814402	BLASTN	1032	1e-77	81
1028	456	700075826H1	SATMON007	g2738247	BLASTN	1038	1e-77	82
1029	456	700104555H1	SATMON010	g886470	BLASTN	1039	1e-77	79
1030	456	700074084H1	SATMON007	g2738247	BLASTN	1039	1e-77	87
1031	456	700076872H1	SATMON007	g1814402	BLASTN	700	1e-76	83
1032	456	700050704H1	SATMON003	g1814402	BLASTN	880	1e-76	82
1033	456	700030215H1	SATMON003	g974781	BLASTN	1019	1e-76	83
1034	456	700077416H1	SATMON007	g886470	BLASTN	1019	1e-76	79
1035	456	700093982H1	SATMON008	g1814402	BLASTN	1019	1e-76	82
1036	456	700048847H1	SATMON003	g2738247	BLASTN	1023	1e-76	85
1037	456	700090739H1	SATMON011	g886470	BLASTN	1026	1e-76	82
1038	456	700092904H1	SATMON008	g886470	BLASTN	1026	1e-76	82

1039	456	LIB3068-009-Q1-K1-E5	LIB3068	g1814402	BLASTN	1027	1e-76	73
1040	456	700336705H1	SATMON019	g1814402	BLASTN	747	1e-75	83
1041	456	700344872H1	SATMON021	g1814402	BLASTN	768	1e-75	85
1042	456	LIB3066-019-Q1-K1-E1	LIB3066	g886470	BLASTN	868	1e-75	81
1043	456	700086322H1	SATMON011	g2738247	BLASTN	1015	1e-75	81
1044	456	LIB3067-052-Q1-K1-B12	LIB3067	g974781	BLASTN	1033	1e-75	81
1045	456	700221395H1	SATMON011	g2738247	BLASTN	1000	1e-74	85
1046	456	700092483H1	SATMON008	g1814402	BLASTN	1001	1e-74	82
1047	456	700618708H1	SATMON034	g886470	BLASTN	1003	1e-74	82
1048	456	700074191H1	SATMON007	g2738247	BLASTN	1005	1e-74	86
1049	456	LIB143-055-Q1-E1-F10	LIB143	g1814402	BLASTN	986	1e-73	83
1050	456	700025955H1	SATMON003	g1814402	BLASTN	993	1e-73	84
1051	456	700574824H1	SATMON030	g886470	BLASTN	564	1e-72	80
1052	456	700201527H1	SATMON003	g974781	BLASTN	812	1e-72	81
1053	456	700092202H1	SATMON008	g886470	BLASTN	946	1e-72	79
1054	456	700223674H1	SATMON011	g2738247	BLASTN	971	1e-72	83
1055	456	700092638H1	SATMON008	g1814402	BLASTN	972	1e-72	84
1056	456	700090025H1	SATMON011	g974781	BLASTN	973	1e-72	81
1057	456	700217069H1	SATMON016	g2738247	BLASTN	976	1e-72	83
1058	456	700349142H1	SATMON023	g1814402	BLASTN	980	1e-72	82
1059	456	700088387H1	SATMON011	g2738247	BLASTN	980	1e-72	85
1060	456	700215102H1	SATMON016	g886470	BLASTN	980	1e-72	79
1061	456	700456708H1	SATMON029	g2738247	BLASTN	982	1e-72	82
1062	456	700210025H1	SATMON016	g1814402	BLASTN	982	1e-72	81
1063	456	700335345H1	SATMON019	g974781	BLASTN	982	1e-72	84
1064	456	700085257H1	SATMON011	g1814402	BLASTN	554	1e-71	82
1065	456	700026781H1	SATMON003	g974781	BLASTN	961	1e-71	82
1066	456	700224265H1	SATMON011	g2738247	BLASTN	962	1e-71	83
1067	456	700083282H1	SATMON011	g1814402	BLASTN	962	1e-71	83
1068	456	700381324H1	SATMON023	g1814402	BLASTN	965	1e-71	82
1069	456	700212017H1	SATMON016	g1814402	BLASTN	640	1e-70	83
1070	456	700088053H1	SATMON011	g886470	BLASTN	853	1e-70	84
1071	456	LIB3062-032-Q1-K1-C2	LIB3062	g1814402	BLASTN	883	1e-70	80
1072	456	700083030H1	SATMON011	g1814402	BLASTN	948	1e-70	81
1073	456	700613980H1	SATMON033	g1814402	BLASTN	950	1e-70	83
1074	456	700347019H1	SATMON021	g974781	BLASTN	952	1e-70	81
1075	456	700335621H1	SATMON019	g1814402	BLASTN	952	1e-70	80
1076	456	700094101H1	SATMON008	g1814402	BLASTN	953	1e-70	83
1077	456	700073128H1	SATMON007	g2738247	BLASTN	954	1e-70	78
1078	456	700076245H1	SATMON007	g1814402	BLASTN	955	1e-70	83
1079	456	700071903H1	SATMON007	g886470	BLASTN	956	1e-70	81
1080	456	700090035H1	SATMON011	g886470	BLASTN	935	1e-69	80
1081	456	700405323H1	SATMON029	g1814402	BLASTN	937	1e-69	83
1082	456	700050121H1	SATMON003	g886470	BLASTN	938	1e-69	83
1083	456	700224714H1	SATMON011	g1814402	BLASTN	939	1e-69	82
1084	456	700281535H2	SATMON019	g1814402	BLASTN	940	1e-69	81
1085	456	700222093H1	SATMON011	g2738247	BLASTN	941	1e-69	81
1086	456	700094659H1	SATMON008	g1814402	BLASTN	942	1e-69	84
1087	456	700237187H1	SATMON010	g1814402	BLASTN	942	1e-69	84
1088	456	700023210H1	SATMON003	g1814402	BLASTN	943	1e-69	83
1089	456	700072813H1	SATMON007	g1814402	BLASTN	944	1e-69	82
1090	456	700075720H1	SATMON007	g1814402	BLASTN	946	1e-69	83
1091	456	700072107H1	SATMON007	g1814402	BLASTN	946	1e-69	80
1092	456	700082225H1	SATMON011	g1814402	BLASTN	691	1e-68	82

1093	456	700085411H1	SATMON011	g1814402	BLASTN	723	1e-68	83
1094	456	700454357H1	SATMON029	g2738247	BLASTN	926	1e-68	82
1095	456	700620976H1	SATMON034	g1814402	BLASTN	927	1e-68	81
1096	456	700241483H1	SATMON010	g1814402	BLASTN	931	1e-68	86
1097	456	700355459H1	SATMON024	g974781	BLASTN	931	1e-68	82
1098	456	700095875H1	SATMON008	g1814402	BLASTN	931	1e-68	87
1099	456	700076702H1	SATMON007	g1814402	BLASTN	934	1e-68	84
1100	456	700212538H1	SATMON016	g1814402	BLASTN	689	1e-67	85
1101	456	700086479H1	SATMON011	g1814402	BLASTN	715	1e-67	82
1102	456	700453684H1	SATMON028	g1814402	BLASTN	747	1e-67	83
1103	456	700204454H1	SATMON003	g1814402	BLASTN	797	1e-67	82
1104	456	700612577H1	SATMON033	g974781	BLASTN	911	1e-67	81
1105	456	700474018H1	SATMON025	g886470	BLASTN	917	1e-67	81
1106	456	700213602H1	SATMON016	g886470	BLASTN	917	1e-67	76
1107	456	700076911H1	SATMON007	g974781	BLASTN	917	1e-67	81
1108	456	700223313H1	SATMON011	g886470	BLASTN	919	1e-67	84
1109	456	700220692H1	SATMON011	g974781	BLASTN	920	1e-67	81
1110	456	700258038H1	SATMON017	g886470	BLASTN	920	1e-67	83
1111	456	700095862H1	SATMON008	g1814402	BLASTN	922	1e-67	86
1112	456	700213396H1	SATMON016	g886470	BLASTN	922	1e-67	83
1113	456	700354259H1	SATMON024	g974781	BLASTN	638	1e-66	81
1114	456	700471505H1	SATMON025	g886470	BLASTN	681	1e-66	83
1115	456	700202439H1	SATMON003	g1814402	BLASTN	706	1e-66	82
1116	456	700223609H1	SATMON011	g974781	BLASTN	770	1e-66	83
1117	456	700405260H1	SATMON028	g1814402	BLASTN	782	1e-66	86
1118	456	700085214H1	SATMON011	g886470	BLASTN	814	1e-66	84
1119	456	700211194H1	SATMON016	g1814402	BLASTN	902	1e-66	84
1120	456	700106641H1	SATMON010	g1814402	BLASTN	903	1e-66	82
1121	456	700405170H1	SATMON028	g974781	BLASTN	904	1e-66	79
1122	456	700090015H1	SATMON011	g886470	BLASTN	905	1e-66	80
1123	456	700623154H1	SATMON034	g886470	BLASTN	906	1e-66	81
1124	456	700458895H1	SATMON029	g886470	BLASTN	906	1e-66	83
1125	456	700094073H1	SATMON008	g886470	BLASTN	907	1e-66	78
1126	456	700158508H1	SATMON012	g2738247	BLASTN	909	1e-66	82
1127	456	700027366H1	SATMON003	g1814402	BLASTN	910	1e-66	81
1128	456	700073636H1	SATMON007	g1814402	BLASTN	615	1e-65	87
1129	456	700220207H1	SATMON011	g886470	BLASTN	785	1e-65	84
1130	456	LIB143-007-Q1-E1-A6	LIB143	g886470	BLASTN	807	1e-65	80
1131	456	700575288H1	SATMON030	g1814402	BLASTN	887	1e-65	83
1132	456	700219870H1	SATMON011	g974781	BLASTN	887	1e-65	82
1133	456	701184573H1	SATMONN06	g974781	BLASTN	888	1e-65	81
1134	456	700076946H1	SATMON007	g1814402	BLASTN	891	1e-65	86
1135	456	700214808H1	SATMON016	g2738247	BLASTN	893	1e-65	81
1136	456	700096274H1	SATMON008	g1814402	BLASTN	894	1e-65	80
1137	456	700160445H1	SATMON012	g1814402	BLASTN	896	1e-65	85
1138	456	700071727H1	SATMON007	g974781	BLASTN	898	1e-65	80
1139	456	700571751H1	SATMON030	g1814402	BLASTN	898	1e-65	78
1140	456	700215107H1	SATMON016	g2738247	BLASTN	507	1e-64	77
1141	456	LIB3067-057-Q1-K1-C6	LIB3067	g2738247	BLASTN	555	1e-64	76
1142	456	700614743H1	SATMON033	g886470	BLASTN	748	1e-64	81
1143	456	700444971H1	SATMON027	g974781	BLASTN	803	1e-64	83
1144	456	700224573H1	SATMON011	g1814402	BLASTN	805	1e-64	86
1145	456	700019276H1	SATMON001	g1814402	BLASTN	875	1e-64	83
1146	456	700210440H1	SATMON016	g2738247	BLASTN	881	1e-64	80

1147	456	700093168H1	SATMON008	g1814402	BLASTN	885	1e-64	86
1148	456	700224478H1	SATMON011	g886470	BLASTN	886	1e-64	81
1149	456	700261958H1	SATMON017	g1814402	BLASTN	781	1e-63	79
1150	456	700074221H1	SATMON007	g1814402	BLASTN	825	1e-63	81
1151	456	700089966H1	SATMON011	g1814402	BLASTN	863	1e-63	84
1152	456	700210422H1	SATMON016	g1814402	BLASTN	863	1e-63	81
1153	456	700160272H1	SATMON012	g1814402	BLASTN	863	1e-63	85
1154	456	700444282H1	SATMON027	g2738247	BLASTN	549	1e-62	84
1155	456	700206489H1	SATMON003	g2738247	BLASTN	597	1e-62	80
1156	456	700446452H1	SATMON027	g974781	BLASTN	711	1e-62	80
1157	456	700265473H1	SATMON017	g1814402	BLASTN	715	1e-62	79
1158	456	700473708H1	SATMON025	g1814402	BLASTN	762	1e-62	82
1159	456	LIB3069-025-Q1-K1-A7	LIB3069	g886470	BLASTN	852	1e-62	80
1160	456	700166631H1	SATMON013	g2738247	BLASTN	858	1e-62	85
1161	456	700582413H1	SATMON031	g1814402	BLASTN	861	1e-62	78
1162	456	700071904H1	SATMON007	g2738247	BLASTN	544	1e-61	77
1163	456	700622655H1	SATMON034	g1814402	BLASTN	592	1e-61	82
1164	456	700082541H1	SATMON011	g1814402	BLASTN	793	1e-61	82
1165	456	700335378H1	SATMON019	g2738247	BLASTN	841	1e-61	76
1166	456	700203585H1	SATMON003	g2738247	BLASTN	846	1e-61	86
1167	456	700053043H1	SATMON007	g2738247	BLASTN	847	1e-61	81
1168	456	700163654H1	SATMON013	g1814402	BLASTN	474	1e-60	84
1169	456	700352269H1	SATMON023	g1814402	BLASTN	479	1e-60	84
1170	456	700106187H1	SATMON010	g1814402	BLASTN	631	1e-60	81
1171	456	LIB143-007-Q1-E1-A7	LIB143	g886470	BLASTN	759	1e-60	76
1172	456	700458694H1	SATMON029	g1814402	BLASTN	827	1e-60	86
1173	456	700213509H1	SATMON016	g886470	BLASTN	834	1e-60	81
1174	456	700150191H1	SATMON007	g2738247	BLASTN	834	1e-60	82
1175	456	700157594H1	SATMON012	g1814402	BLASTN	835	1e-60	80
1176	456	700094046H1	SATMON008	g974781	BLASTN	836	1e-60	79
1177	456	700220254H1	SATMON011	g2738247	BLASTN	687	1e-59	77
1178	456	700095485H1	SATMON008	g886470	BLASTN	819	1e-59	77
1179	456	LIB143-006-Q1-E1-B3	LIB143	g1814402	BLASTN	830	1e-59	65
1180	456	700071763H1	SATMON007	g2738247	BLASTN	565	1e-58	75
1181	456	700082259H1	SATMON011	g1814402	BLASTN	707	1e-58	84
1182	456	700020717H1	SATMON001	g886470	BLASTN	805	1e-58	81
1183	456	700041892H1	SATMON004	g1814402	BLASTN	805	1e-58	86
1184	456	700356573H1	SATMON024	g1814402	BLASTN	811	1e-58	81
1185	456	700171632H1	SATMON013	g2738247	BLASTN	811	1e-58	85
1186	456	700222516H1	SATMON011	g1814402	BLASTN	812	1e-58	87
1187	456	700466679H1	SATMON025	g1814402	BLASTN	812	1e-58	87
1188	456	700208929H1	SATMON016	g1814402	BLASTN	407	1e-57	78
1189	456	700355014H1	SATMON024	g1814402	BLASTN	592	1e-57	85
1190	456	701159834H1	SATMONN04	g1814402	BLASTN	683	1e-57	77
1191	456	700351137H1	SATMON023	g974781	BLASTN	725	1e-57	79
1192	456	700027308H1	SATMON003	g2738247	BLASTN	791	1e-57	80
1193	456	700142587H1	SATMON012	g974781	BLASTN	791	1e-57	83
1194	456	700170157H1	SATMON013	g1814402	BLASTN	791	1e-57	81
1195	456	700019438H1	SATMON001	g974781	BLASTN	798	1e-57	82
1196	456	700166430H1	SATMON013	g886470	BLASTN	799	1e-57	84
1197	456	700047369H1	SATMON003	g886470	BLASTN	799	1e-57	80
1198	456	700152373H1	SATMON007	g2738247	BLASTN	799	1e-57	84
1199	456	700050824H1	SATMON003	g2738247	BLASTN	605	1e-56	81
1200	456	700204588H1	SATMON003	g886470	BLASTN	720	1e-56	82

1201	456	700071667H1	SATMON007	g1814402	BLASTN	787	1e-56	85
1202	456	701185269H1	SATMONN06	g886470	BLASTN	598	1e-55	76
1203	456	700243844H1	SATMON010	g2738247	BLASTN	769	1e-55	79
1204	456	700221969H1	SATMON011	g974781	BLASTN	771	1e-55	79
1205	456	700335706H1	SATMON019	g1814402	BLASTN	773	1e-55	80
1206	456	700622285H1	SATMON034	g1814402	BLASTN	774	1e-55	82
1207	456	700089665H1	SATMON011	g2738247	BLASTN	774	1e-55	78
1208	456	700464833H1	SATMON025	g2738247	BLASTN	455	1e-54	72
1209	456	LIB3069-057-Q1-K1-A6	LIB3069	g974781	BLASTN	564	1e-54	76
1210	456	700156818H1	SATMON012	g974781	BLASTN	757	1e-54	81
1211	456	700219420H1	SATMON011	g974781	BLASTN	758	1e-54	79
1212	456	700152570H1	SATMON007	g1814402	BLASTN	761	1e-54	86
1213	456	700084484H1	SATMON011	g1814402	BLASTN	764	1e-54	84
1214	456	700152208H1	SATMON007	g1814402	BLASTN	765	1e-54	82
1215	456	700550889H1	SATMON022	g2738247	BLASTN	546	1e-53	76
1216	456	700025869H1	SATMON003	g2738247	BLASTN	675	1e-53	79
1217	456	700383075H1	SATMON024	g974781	BLASTN	724	1e-53	81
1218	456	700575104H1	SATMON030	g1814402	BLASTN	745	1e-53	70
1219	456	LIB3069-022-Q1-K1-C1	LIB3069	g974781	BLASTN	768	1e-53	81
1220	456	700215272H1	SATMON016	g1814402	BLASTN	674	1e-52	83
1221	456	700167881H1	SATMON013	g974781	BLASTN	737	1e-52	79
1222	456	700019727H1	SATMON001	g2738247	BLASTN	737	1e-52	81
1223	456	700351776H1	SATMON023	g886470	BLASTN	741	1e-52	76
1224	456	700210543H1	SATMON016	g974781	BLASTN	742	1e-52	85
1225	456	700571604H1	SATMON030	g1814402	BLASTN	360	1e-51	76
1226	456	700169777H1	SATMON013	g1814402	BLASTN	426	1e-51	83
1227	456	700096744H1	SATMON008	g2738247	BLASTN	539	1e-51	79
1228	456	700383157H1	SATMON024	g974781	BLASTN	667	1e-51	79
1229	456	700155764H1	SATMON007	g886470	BLASTN	721	1e-51	75
1230	456	700150817H1	SATMON007	g1814402	BLASTN	727	1e-51	75
1231	456	700619660H1	SATMON034	g886470	BLASTN	727	1e-51	73
1232	456	700471085H1	SATMON025	g1814402	BLASTN	406	1e-50	79
1233	456	700163226H1	SATMON013	g2738247	BLASTN	475	1e-50	78
1234	456	700444567H1	SATMON027	g1814402	BLASTN	498	1e-50	81
1235	456	700457303H1	SATMON029	g2738247	BLASTN	662	1e-50	80
1236	456	700088065H1	SATMON011	g886470	BLASTN	707	1e-50	79
1237	456	700152592H1	SATMON007	g886470	BLASTN	716	1e-50	80
1238	456	LIB3068-061-Q1-K1-B2	LIB3068	g886470	BLASTN	730	1e-50	78
1239	456	700331880H1	SATMON019	g1814402	BLASTN	697	1e-49	85
1240	456	700218025H1	SATMON016	g886470	BLASTN	700	1e-49	79
1241	456	700348643H1	SATMON023	g1814402	BLASTN	701	1e-49	80
1242	456	700236975H1	SATMON010	g974781	BLASTN	704	1e-49	81
1243	456	700442968H1	SATMON026	g886470	BLASTN	418	1e-48	79
1244	456	700479529H1	SATMON034	g2738247	BLASTN	508	1e-48	75
1245	456	700160983H1	SATMON012	g1814402	BLASTN	690	1e-48	83
1246	456	700050185H1	SATMON003	g974781	BLASTN	421	1e-47	75
1247	456	700611764H1	SATMON022	g1814402	BLASTN	503	1e-47	79
1248	456	700622669H1	SATMON034	g1814402	BLASTN	549	1e-47	79
1249	456	700153237H1	SATMON007	g1814402	BLASTN	672	1e-47	80
1250	456	700242703H1	SATMON010	g886470	BLASTN	682	1e-47	81
1251	456	700165177H1	SATMON013	g2738247	BLASTN	682	1e-47	78
1252	456	700379889H1	SATMON021	g886470	BLASTN	682	1e-47	85
1253	456	700449243H1	SATMON028	g1814402	BLASTN	357	1e-46	83
1254	456	700453282H1	SATMON028	g886470	BLASTN	608	1e-46	81

1255	456	700224308H1	SATMON011	g1814402	BLASTN	661	1e-46	83
1256	456	700150484H1	SATMON007	g886470	BLASTN	663	1e-46	78
1257	456	700240609H1	SATMON010	g2738247	BLASTN	668	1e-46	79
1258	456	700171610H1	SATMON013	g974781	BLASTN	669	1e-46	79
1259	456	LIB143-027-Q1-E1-H3	LIB143	g1814402	BLASTN	670	1e-46	86
1260	456	700334084H1	SATMON019	g1814402	BLASTN	467	1e-45	81
1261	456	LIB3069-043-Q1-K1-H6	LIB3069	g886470	BLASTN	546	1e-45	83
1262	456	700222515H1	SATMON011	g1814402	BLASTN	648	1e-45	75
1263	456	700223749H1	SATMON011	g1814402	BLASTN	648	1e-45	75
1264	456	701180187H1	SATMONN05	g974781	BLASTN	648	1e-45	78
1265	456	700050111H1	SATMON003	g2738247	BLASTN	653	1e-45	82
1266	456	700051275H1	SATMON003	g1814402	BLASTN	379	1e-44	75
1267	456	700235202H1	SATMON010	g2738247	BLASTN	637	1e-44	80
1268	456	700257385H1	SATMON017	g1814402	BLASTN	638	1e-44	80
1269	456	LIB143-028-Q1-E1-H7	LIB143	g1814402	BLASTN	638	1e-44	80
1270	456	700576066H1	SATMON030	g886470	BLASTN	516	1e-43	76
1271	456	700455065H1	SATMON029	g886470	BLASTN	624	1e-43	79
1272	456	700074158H1	SATMON007	g1814402	BLASTN	630	1e-43	83
1273	456	700171322H1	SATMON013	g1814402	BLASTN	634	1e-43	77
1274	456	700457369H1	SATMON029	g974781	BLASTN	392	1e-42	83
1275	456	700378252H1	SATMON019	g2738247	BLASTN	462	1e-42	77
1276	456	700454626H1	SATMON029	g1814402	BLASTN	495	1e-42	85
1277	456	700208182H1	SATMON016	g2738247	BLASTN	564	1e-42	73
1278	456	700618848H1	SATMON034	g886470	BLASTN	615	1e-42	80
1279	456	700222969H1	SATMON011	g1814402	BLASTN	622	1e-42	74
1280	456	700444782H1	SATMON027	g1814402	BLASTN	337	1e-41	81
1281	456	700204404H1	SATMON003	g2738247	BLASTN	607	1e-41	80
1282	456	700347028H1	SATMON021	g1814402	BLASTN	598	1e-40	81
1283	456	700172447H1	SATMON013	g1814402	BLASTN	328	1e-39	85
1284	456	700549696H1	SATMON022	g886470	BLASTN	360	1e-39	81
1285	456	700257287H1	SATMON017	g1814402	BLASTN	365	1e-38	84
1286	456	700569630H1	SATMON030	g1814402	BLASTN	566	1e-38	78
1287	456	700207258H1	SATMON017	g1814402	BLASTN	568	1e-38	85
1288	456	700052467H1	SATMON003	g1814402	BLASTN	515	1e-37	82
1289	456	700083656H1	SATMON011	g2738247	BLASTN	552	1e-37	83
1290	456	700429279H1	SATMONN01	g2738247	BLASTN	556	1e-37	79
1291	456	700349921H1	SATMON023	g2738247	BLASTN	471	1e-36	78
1292	456	700449609H1	SATMON028	g886470	BLASTN	540	1e-36	81
1293	456	700150152H1	SATMON007	g1814402	BLASTN	543	1e-36	74
1294	456	700075675H1	SATMON007	g1814402	BLASTN	546	1e-36	81
1295	456	700267015H1	SATMON017	g886470	BLASTN	550	1e-36	75
1296	456	700465118H1	SATMON025	g1814402	BLASTN	555	1e-36	78
1297	456	700456662H1	SATMON029	g2738247	BLASTN	314	1e-34	76
1298	456	700405315H1	SATMON029	g886470	BLASTN	519	1e-34	83
1299	456	700218639H1	SATMON011	g974781	BLASTN	524	1e-34	81
1300	456	700216606H1	SATMON016	g1814402	BLASTN	524	1e-34	83
1301	456	700456965H1	SATMON029	g2738247	BLASTN	525	1e-34	78
1302	456	700102147H1	SATMON010	g974781	BLASTN	526	1e-34	82
1303	456	700235734H1	SATMON010	g2738247	BLASTN	526	1e-34	83
1304	456	LIB3068-009-Q1-K1-E10	LIB3068	g886470	BLASTN	548	1e-34	70
1305	456	700029363H1	SATMON003	g974781	BLASTN	377	1e-33	80
1306	456	LIB3069-030-Q1-K1-D10	LIB3069	g886470	BLASTN	297	1e-30	78
1307	456	700334895H1	SATMON019	g974781	BLASTN	468	1e-30	82
1308	456	700150963H1	SATMON007	g974781	BLASTN	459	1e-29	88

1309	456	700152817H1	SATMON007	g974781	BLASTN	459	1e-29	88
1310	456	700208732H1	SATMON016	g974781	BLASTN	485	1e-29	76
1311	456	700051461H1	SATMON003	g886470	BLASTN	467	1e-28	73
1312	456	700616589H1	SATMON033	g2738247	BLASTN	273	1e-27	78
1313	456	700236150H1	SATMON010	g974781	BLASTN	434	1e-27	84
1314	456	700453369H1	SATMON028	g1814402	BLASTN	436	1e-27	84
1315	456	700621739H1	SATMON034	g1814402	BLASTN	436	1e-27	69
1316	456	700075162H1	SATMON007	g1814402	BLASTN	438	1e-27	85
1317	456	700405040H1	SATMON027	g974781	BLASTN	438	1e-27	72
1318	456	700356893H1	SATMON024	g974781	BLASTN	458	1e-27	82
1319	456	701185493H1	SATMONN06	g886471	BLASTX	72	1e-26	97
1320	456	700206095H1	SATMON003	g974781	BLASTN	421	1e-26	83
1321	456	700083901H1	SATMON011	g1814402	BLASTN	421	1e-26	80
1322	456	LIB3062-026-Q1-K1-D7	LIB3062	g1814402	BLASTN	423	1e-24	81
1323	456	700377277H1	SATMON019	g1814402	BLASTN	357	1e-20	75
1324	456	700201214H1	SATMON003	g2738248	BLASTX	152	1e-18	77
1325	456	700025959H1	SATMON003	g886471	BLASTX	188	1e-18	94
1326	456	700259484H1	SATMON017	g1814402	BLASTN	189	1e-17	82
1327	456	700613969H1	SATMON033	g886470	BLASTN	315	1e-17	90
1328	456	700215602H1	SATMON016	g2738248	BLASTX	158	1e-16	74
1329	456	700438152H1	SATMON026	g2738247	BLASTN	258	1e-16	75
1330	456	700458654H1	SATMON029	g886470	BLASTN	330	1e-16	70
1331	456	700449563H1	SATMON028	g886471	BLASTX	163	1e-15	89
1332	456	700236490H1	SATMON010	g1814402	BLASTN	276	1e-14	89
1333	456	700456927H1	SATMON029	g1814403	BLASTX	119	1e-13	86
1334	456	701180088H1	SATMONN05	g886471	BLASTX	152	1e-13	60
1335	456	700455716H1	SATMON029	g2738248	BLASTX	111	1e-12	67
1336	456	700573520H1	SATMON030	g886471	BLASTX	82	1e-11	84
1337	456	700569938H1	SATMON030	g2738248	BLASTX	132	1e-11	91
1338	456	700551958H1	SATMON022	g2738247	BLASTN	264	1e-11	82
1339	456	700337475H1	SATMON020	g2738247	BLASTN	268	1e-11	83
1340	456	700170827H1	SATMON013	g886471	BLASTX	124	1e-10	77
1341	456	700453382H1	SATMON028	g974781	BLASTN	152	1e-10	85
1342	456	700089211H1	SATMON011	g2738247	BLASTN	245	1e-9	79
1343	456	700103155H1	SATMON010	g2738248	BLASTX	82	1e-8	85
1344	5523	LIB3062-017-Q1-K1-F4	LIB3062	g2738247	BLASTN	932	1e-84	76
1345	5523	700210708H1	SATMON016	g2738247	BLASTN	952	1e-70	79
1346	5523	700219307H1	SATMON011	g1814402	BLASTN	892	1e-65	78
1347	5523	700221188H1	SATMON011	g2738247	BLASTN	867	1e-63	81
1348	5523	700203884H1	SATMON003	g2738247	BLASTN	874	1e-63	77
1349	5523	700218549H1	SATMON011	g2738247	BLASTN	811	1e-58	77
1350	5523	700572845H2	SATMON030	g2738247	BLASTN	785	1e-56	77
1351	5523	700152362H1	SATMON007	g886470	BLASTN	742	1e-52	79
1352	5523	700152138H1	SATMON007	g2738247	BLASTN	714	1e-50	80
1353	5523	700218842H1	SATMON011	g2738247	BLASTN	700	1e-49	75
2666	-700697958	700697958H1	SOYMON015	g2738247	BLASTN	258	1e-10	83
2667	-700731277	700731277H1	SOYMON009	g886470	BLASTN	930	1e-68	83
2668	-700749261	700749261H1	SOYMON013	g1814402	BLASTN	508	1e-38	78
2669	-700787742	700787742H2	SOYMON011	g1814402	BLASTN	648	1e-50	81
2670	-700831146	700831146H1	SOYMON019	g1814402	BLASTN	476	1e-34	80
2671	-700832029	700832029H1	SOYMON019	g1814402	BLASTN	328	1e-27	78
2672	-700854481	700854481H1	SOYMON023	g1814402	BLASTN	193	1e-11	85
2673	-700873716	700873716H1	SOYMON018	g1749542	BLASTX	99	1e-15	53
2674	-700893904	700893904H1	SOYMON024	g886470	BLASTN	413	1e-25	82

2675	-700909658	700909658H1	SOYMON022	g1814402	BLASTN	251	1e-10	91
2676	-700943841	700943841H1	SOYMON024	g886470	BLASTN	549	1e-49	82
2677	-700963223	700963223H1	SOYMON022	g886470	BLASTN	649	1e-45	72
2678	-700974103	700974103H1	SOYMON005	g886470	BLASTN	763	1e-54	80
2679	-700994293	700994293H1	SOYMON011	g974782	BLASTX	97	1e-12	72
2680	-701004816	701004816H1	SOYMON019	g1814402	BLASTN	582	1e-39	84
2681	-701007125	701007125H1	SOYMON019	g1814403	BLASTX	152	1e-13	87
2682	-701008540	701008540H1	SOYMON019	g886470	BLASTN	770	1e-55	74
2683	-701037766	701037766H1	SOYMON029	g1814403	BLASTX	72	1e-10	69
2684	-701062191	701062191H1	SOYMON033	g974781	BLASTN	225	1e-9	79
2685	-701105474	701105474H1	SOYMON036	g974782	BLASTX	164	1e-15	91
2686	-GM14442	LIB3049-056-Q1-E1-B1	LIB3049	g974781	BLASTN	231	1e-10	79
2687	-GM19631	LIB3056-007-Q1-N1-G9	LIB3056	g974781	BLASTN	388	1e-37	78
2688	-GM37189	LIB3051-072-Q1-K1-B5	LIB3051	g1814402	BLASTN	316	1e-15	78
2689	-GM44802	LIB3053-004-Q1-N1-B2	LIB3053	g974782	BLASTX	85	1e-26	90
2690	1382	700683236H1	SOYMON008	g886470	BLASTN	805	1e-58	78
2691	1382	700566434H1	SOYMON002	g886471	BLASTX	162	1e-23	76
2692	15690	701064361H1	SOYMON034	g974781	BLASTN	951	1e-70	84
2693	15690	700847301H1	SOYMON021	g886470	BLASTN	707	1e-66	85
2694	17335	LIB3051-088-Q1-K1-D9	LIB3051	g886470	BLASTN	1285	1e-101	79
2695	17335	701003832H1	SOYMON019	g974781	BLASTN	811	1e-58	77
2696	17335	700864888H1	SOYMON016	g1814402	BLASTN	767	1e-55	79
2697	17335	700672491H1	SOYMON006	g974781	BLASTN	428	1e-46	75
2698	17335	701003457H1	SOYMON019	g1814402	BLASTN	411	1e-29	84
2699	17335	700833672H1	SOYMON019	g974781	BLASTN	445	1e-28	79
2700	17900	700850578H1	SOYMON023	g2738247	BLASTN	905	1e-66	82
2701	17900	701053154H1	SOYMON032	g1814402	BLASTN	725	1e-65	83
2702	17900	700842351H1	SOYMON020	g1814402	BLASTN	878	1e-64	83
2703	17900	700837656H1	SOYMON020	g1814402	BLASTN	841	1e-61	83
2704	17900	700890851H1	SOYMON024	g974781	BLASTN	747	1e-53	80
2705	20688	700908840H1	SOYMON022	g2738247	BLASTN	889	1e-65	82
2706	20688	700908848H1	SOYMON022	g2738247	BLASTN	877	1e-64	81
2707	33542	LIB3051-009-Q1-E1-E5	LIB3051	g974781	BLASTN	1218	1e-92	79
2708	33542	700748773H1	SOYMON013	g974781	BLASTN	669	1e-46	78
2709	33542	700836363H1	SOYMON020	g2738247	BLASTN	596	1e-40	80
2710	4243	701123616H1	SOYMON037	g1814402	BLASTN	995	1e-74	87
2711	4243	700555001H1	SOYMON001	g1814402	BLASTN	975	1e-72	90
2712	4243	701002967H1	SOYMON019	g1814402	BLASTN	950	1e-70	86
2713	4243	700653509H1	SOYMON003	g1814402	BLASTN	539	1e-69	85
2714	4243	701206028H1	SOYMON035	g1814402	BLASTN	938	1e-69	86
2715	4243	700962115H1	SOYMON022	g1814402	BLASTN	923	1e-68	86
2716	4243	700866243H1	SOYMON016	g1814402	BLASTN	909	1e-66	82
2717	4243	700752507H1	SOYMON014	g1814402	BLASTN	887	1e-65	85
2718	4243	701003887H1	SOYMON019	g1814402	BLASTN	863	1e-63	86
2719	4243	700556913H1	SOYMON001	g1814402	BLASTN	865	1e-63	86
2720	4243	701013549H1	SOYMON019	g1814402	BLASTN	867	1e-63	90
2721	4243	701209706H1	SOYMON035	g1814402	BLASTN	871	1e-63	90
2722	4243	701010487H1	SOYMON019	g1814402	BLASTN	529	1e-62	79
2723	4243	700548246H1	SOYMON002	g1814402	BLASTN	553	1e-62	82
2724	4243	701138219H1	SOYMON038	g1814402	BLASTN	594	1e-62	86
2725	4243	700965160H1	SOYMON022	g1814402	BLASTN	852	1e-62	91
2726	4243	701015168H1	SOYMON019	g1814402	BLASTN	855	1e-62	89
2727	4243	701136095H1	SOYMON038	g1814402	BLASTN	839	1e-61	88
2728	4243	700761789H1	SOYMON015	g1814402	BLASTN	845	1e-61	86

2729	4243	701105695H1	SOYMON036	g1814402	BLASTN	835	1e-60	87
2730	4243	700991714H1	SOYMON011	g1814402	BLASTN	502	1e-59	83
2731	4243	700564223H1	SOYMON002	g1814402	BLASTN	557	1e-59	90
2732	4243	700987384H1	SOYMON009	g1814402	BLASTN	803	1e-58	86
2733	4243	700833934H1	SOYMON019	g1814402	BLASTN	806	1e-58	89
2734	4243	700835181H1	SOYMON019	g1814402	BLASTN	806	1e-58	82
2735	4243	700737529H1	SOYMON010	g1814402	BLASTN	810	1e-58	92
2736	4243	701012851H1	SOYMON019	g1814402	BLASTN	811	1e-58	91
2737	4243	700556592H1	SOYMON001	g1814402	BLASTN	814	1e-58	88
2738	4243	700907579H1	SOYMON022	g1814402	BLASTN	781	1e-56	89
2739	4243	700961749H1	SOYMON022	g1814402	BLASTN	785	1e-56	91
2740	4243	700835239H1	SOYMON019	g1814402	BLASTN	787	1e-56	86
2741	4243	700646425H1	SOYMON013	g1814402	BLASTN	772	1e-55	89
2742	4243	701123924H1	SOYMON037	g1814402	BLASTN	775	1e-55	91
2743	4243	700957759H1	SOYMON022	g1814402	BLASTN	776	1e-55	90
2744	4243	700964425H1	SOYMON022	g1814402	BLASTN	777	1e-55	86
2745	4243	700962173H1	SOYMON022	g1814402	BLASTN	778	1e-55	91
2746	4243	701066293H1	SOYMON034	g1814402	BLASTN	759	1e-54	81
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2750	4243	701060510H1	SOYMON033	g1814402	BLASTN	748	1e-53	84
2751	4243	700848730H1	SOYMON021	g1814402	BLASTN	749	1e-53	90
2752	4243	700754870H1	SOYMON014	g1814402	BLASTN	751	1e-53	84
2753	4243	701212482H1	SOYMON035	g1814402	BLASTN	751	1e-53	84
2754	4243	700753283H1	SOYMON014	g1814402	BLASTN	752	1e-53	86
2755	4243	700738517H1	SOYMON012	g1814402	BLASTN	636	1e-52	89
2756	4243	700833978H1	SOYMON019	g1814402	BLASTN	740	1e-52	91
2757	4243	700756424H1	SOYMON014	g1814402	BLASTN	729	1e-51	82
2758	4243	701011759H1	SOYMON019	g1814402	BLASTN	729	1e-51	91
2759	4243	701010103H2	SOYMON019	g1814402	BLASTN	707	1e-50	82
2760	4243	700741392H1	SOYMON012	g1814402	BLASTN	707	1e-50	84
2761	4243	701123062H1	SOYMON037	g1814402	BLASTN	308	1e-49	88
2762	4243	701048949H1	SOYMON032	g1814402	BLASTN	502	1e-49	85
2763	4243	700834566H1	SOYMON019	g1814402	BLASTN	618	1e-49	88
2764	4243	700963965H1	SOYMON022	g1814402	BLASTN	685	1e-48	78
2765	4243	700986376H1	SOYMON009	g1814402	BLASTN	694	1e-48	84
2766	4243	701012708H1	SOYMON019	g1814402	BLASTN	521	1e-47	91
2767	4243	700746927H1	SOYMON013	g1814402	BLASTN	547	1e-46	78
2768	4243	700997448H1	SOYMON018	g1814402	BLASTN	470	1e-45	89
2769	4243	700830667H1	SOYMON019	g1814402	BLASTN	647	1e-45	86
2770	4243	700891479H1	SOYMON024	g1814402	BLASTN	654	1e-45	88
2771	4243	700562246H1	SOYMON002	g1814402	BLASTN	656	1e-45	86
2772	4243	701097325H1	SOYMON028	g1814402	BLASTN	420	1e-44	92
2773	4243	700835830H1	SOYMON019	g1814402	BLASTN	503	1e-44	86
2774	4243	700962969H1	SOYMON022	g1814402	BLASTN	515	1e-44	89
2775	4243	701102860H1	SOYMON028	g1814402	BLASTN	309	1e-42	92
2776	4243	700763506H1	SOYMON015	g1814402	BLASTN	456	1e-42	84
2777	4243	700994059H1	SOYMON011	g1814402	BLASTN	487	1e-42	87
2778	4243	700751551H1	SOYMON014	g1814402	BLASTN	568	1e-41	85
2779	4243	701108872H1	SOYMON036	g1814402	BLASTN	603	1e-41	85
2780	4243	700994053H1	SOYMON011	g1814402	BLASTN	587	1e-40	85
2781	4243	700729064H1	SOYMON009	g1814402	BLASTN	413	1e-38	90
2782	4243	700874176H1	SOYMON018	g1814402	BLASTN	468	1e-38	92

2783	4243	700742963H1	SOYMON012	g1814402	BLASTN	565	1e-38	86
2784	4243	701212923H1	SOYMON035	g1814402	BLASTN	572	1e-38	84
2785	4243	700994851H1	SOYMON011	g1814402	BLASTN	548	1e-36	76
2786	4243	701000193H1	SOYMON018	g1814402	BLASTN	296	1e-33	87
2787	4243	700756219H1	SOYMON014	g1814402	BLASTN	394	1e-32	80
2788	4243	701014751H1	SOYMON019	g1814402	BLASTN	461	1e-29	90
2789	4243	700561163H1	SOYMON002	g1814402	BLASTN	231	1e-25	89
2790	4243	700650284H1	SOYMON003	g974782	BLASTX	157	1e-14	96
2791	4243	700869218H1	SOYMON016	g1814402	BLASTN	248	1e-11	93
2792	550	LIB3028-007-Q1-B1-B6	LIB3028	g886470	BLASTN	1440	1e-111	81
2793	550	LIB3040-017-Q1-E1-E8	LIB3040	g886470	BLASTN	1260	1e-96	80
2794	550	700650656H1	SOYMON003	g1814402	BLASTN	1072	1e-93	83
2795	550	LIB3051-091-Q1-K1-C2	LIB3051	g886470	BLASTN	1124	1e-89	80
2796	550	LIB3051-072-Q1-K1-B3	LIB3051	g974781	BLASTN	1114	1e-83	82
2797	550	LIB3051-006-Q1-E1-G9	LIB3051	g974781	BLASTN	703	1e-82	81
2798	550	700563811H1	SOYMON002	g1814402	BLASTN	1071	1e-80	84
2799	550	700754104H1	SOYMON014	g974781	BLASTN	1052	1e-78	86
2800	550	701002973H1	SOYMON019	g1814402	BLASTN	1039	1e-77	85
2801	550	700986733H1	SOYMON009	g974781	BLASTN	1039	1e-77	83
2802	550	700557595H1	SOYMON001	g886470	BLASTN	1019	1e-76	83
2803	550	700563477H1	SOYMON002	g2738247	BLASTN	1024	1e-76	85
2804	550	700976112H1	SOYMON009	g886470	BLASTN	1024	1e-76	84
2805	550	701004484H1	SOYMON019	g1814402	BLASTN	1007	1e-75	84
2806	550	700889161H1	SOYMON024	g974781	BLASTN	996	1e-74	85
2807	550	700833110H1	SOYMON019	g974781	BLASTN	998	1e-74	87
2808	550	701124559H1	SOYMON037	g974781	BLASTN	1002	1e-74	86
2809	550	700729207H1	SOYMON009	g974781	BLASTN	1004	1e-74	85
2810	550	700730012H1	SOYMON009	g886470	BLASTN	987	1e-73	85
2811	550	700987128H1	SOYMON009	g2738247	BLASTN	989	1e-73	85
2812	550	700564788H1	SOYMON002	g974781	BLASTN	990	1e-73	84
2813	550	701099104H1	SOYMON028	g1814402	BLASTN	994	1e-73	86
2814	550	LIB3065-008-Q1-N1-B4	LIB3065	g2738247	BLASTN	839	1e-72	83
2815	550	701104283H1	SOYMON036	g974781	BLASTN	976	1e-72	83
2816	550	700726387H1	SOYMON009	g974781	BLASTN	980	1e-72	84
2817	550	700900884H1	SOYMON027	g1814402	BLASTN	982	1e-72	85
2818	550	LIB3051-029-Q1-K1-D6	LIB3051	g886470	BLASTN	824	1e-71	81
2819	550	701213995H1	SOYMON035	g1814402	BLASTN	962	1e-71	84
2820	550	700683596H1	SOYMON008	g1814402	BLASTN	968	1e-71	85
2821	550	700646529H1	SOYMON014	g1814402	BLASTN	867	1e-70	84
2822	550	700756704H1	SOYMON014	g1814402	BLASTN	893	1e-70	87
2823	550	700991647H1	SOYMON011	g1814402	BLASTN	947	1e-70	84
2824	550	700962195H1	SOYMON022	g974781	BLASTN	947	1e-70	87
2825	550	700994369H1	SOYMON011	g1814402	BLASTN	949	1e-70	84
2826	550	700672477H1	SOYMON006	g1814402	BLASTN	951	1e-70	85
2827	550	700946215H1	SOYMON024	g2738247	BLASTN	957	1e-70	83
2828	550	701152765H1	SOYMON031	g974781	BLASTN	958	1e-70	88
2829	550	701010552H1	SOYMON019	g974781	BLASTN	501	1e-69	83
2830	550	LIB3056-004-Q1-N1-B12	LIB3056	g2738247	BLASTN	698	1e-69	77
2831	550	701100656H1	SOYMON028	g974781	BLASTN	890	1e-69	85
2832	550	700985362H1	SOYMON009	g1814402	BLASTN	937	1e-69	82
2833	550	700746178H1	SOYMON013	g974781	BLASTN	938	1e-69	85
2834	550	700674440H1	SOYMON007	g974781	BLASTN	938	1e-69	83
2835	550	700556934H1	SOYMON001	g1814402	BLASTN	938	1e-69	82
2836	550	700652624H1	SOYMON003	g1814402	BLASTN	939	1e-69	82

2837	550	700952331H1	SOYMON022	g886470	BLASTN	946	1e-69	83
2838	550	700725576H1	SOYMON009	g886470	BLASTN	946	1e-69	85
2839	550	701003602H1	SOYMON019	g2738247	BLASTN	924	1e-68	83
2840	550	700745053H1	SOYMON013	g974781	BLASTN	924	1e-68	85
2841	550	700895781H1	SOYMON027	g1814402	BLASTN	926	1e-68	84
2842	550	700664593H1	SOYMON005	g974781	BLASTN	926	1e-68	87
2843	550	700864264H1	SOYMON016	g1814402	BLASTN	930	1e-68	84
2844	550	700674466H1	SOYMON007	g974781	BLASTN	933	1e-68	83
2845	550	700564170H1	SOYMON002	g974781	BLASTN	861	1e-67	83
2846	550	700654531H1	SOYMON004	g974781	BLASTN	911	1e-67	81
2847	550	700996341H1	SOYMON018	g886470	BLASTN	911	1e-67	83
2848	550	701136257H1	SOYMON038	g886470	BLASTN	912	1e-67	82
2849	550	700751114H1	SOYMON014	g1814402	BLASTN	914	1e-67	84
2850	550	700657606H1	SOYMON004	g1814402	BLASTN	916	1e-67	87
2851	550	700983827H1	SOYMON009	g974781	BLASTN	918	1e-67	85
2852	550	700981249H1	SOYMON009	g1814402	BLASTN	921	1e-67	80
2853	550	700836103H1	SOYMON019	g886470	BLASTN	922	1e-67	82
2854	550	701011869H1	SOYMON019	g886470	BLASTN	486	1e-66	86
2855	550	701097045H1	SOYMON028	g886470	BLASTN	601	1e-66	85
2856	550	701012695H1	SOYMON019	g974781	BLASTN	777	1e-66	85
2857	550	700945396H1	SOYMON024	g974781	BLASTN	902	1e-66	85
2858	550	700755390H1	SOYMON014	g974781	BLASTN	903	1e-66	86
2859	550	700967721H1	SOYMON033	g886470	BLASTN	910	1e-66	82
2860	550	700750610H1	SOYMON014	g974781	BLASTN	910	1e-66	84
2861	550	700908231H1	SOYMON022	g974781	BLASTN	527	1e-65	83
2862	550	701003835H1	SOYMON019	g886470	BLASTN	723	1e-65	82
2863	550	700790802H1	SOYMON011	g2738247	BLASTN	891	1e-65	82
2864	550	701053438H1	SOYMON032	g974781	BLASTN	893	1e-65	82
2865	550	701009430H1	SOYMON019	g974781	BLASTN	894	1e-65	85
2866	550	700891415H1	SOYMON024	g1814402	BLASTN	895	1e-65	84
2867	550	701100681H1	SOYMON028	g2738247	BLASTN	482	1e-64	84
2868	550	700893420H1	SOYMON024	g974781	BLASTN	570	1e-64	89
2869	550	700752741H1	SOYMON014	g886470	BLASTN	880	1e-64	82
2870	550	700955938H1	SOYMON022	g1814402	BLASTN	882	1e-64	81
2871	550	701015042H1	SOYMON019	g2738247	BLASTN	886	1e-64	82
2872	550	LIB3051-072-Q1-K1-B1	LIB3051	g974781	BLASTN	645	1e-63	76
2873	550	700741918H1	SOYMON012	g886470	BLASTN	722	1e-63	84
2874	550	701103319H1	SOYMON028	g974781	BLASTN	792	1e-63	82
2875	550	700833218H1	SOYMON019	g2738247	BLASTN	863	1e-63	81
2876	550	701008071H1	SOYMON019	g886470	BLASTN	867	1e-63	83
2877	550	700832073H1	SOYMON019	g974781	BLASTN	871	1e-63	83
2878	550	700889695H1	SOYMON024	g974781	BLASTN	872	1e-63	83
2879	550	701007489H2	SOYMON019	g974781	BLASTN	873	1e-63	84
2880	550	700895858H1	SOYMON027	g974781	BLASTN	874	1e-63	84
2881	550	700753955H1	SOYMON014	g974781	BLASTN	470	1e-62	87
2882	550	700564433H1	SOYMON002	g886470	BLASTN	757	1e-62	84
2883	550	700963115H1	SOYMON022	g974781	BLASTN	851	1e-62	83
2884	550	700894728H1	SOYMON024	g886470	BLASTN	860	1e-62	84
2885	550	701056915H1	SOYMON033	g886470	BLASTN	862	1e-62	84
2886	550	700741134H1	SOYMON012	g886470	BLASTN	862	1e-62	83
2887	550	700847591H1	SOYMON021	g974781	BLASTN	842	1e-61	84
2888	550	700941253H1	SOYMON024	g2738247	BLASTN	844	1e-61	80
2889	550	701004315H1	SOYMON019	g2738247	BLASTN	846	1e-61	83
2890	550	700895720H1	SOYMON027	g974781	BLASTN	848	1e-61	83

2891	550	701013541H1	SOYMON019	g974781	BLASTN	849	1e-61	84
2892	550	700892552H1	SOYMON024	g886470	BLASTN	719	1e-60	83
2893	550	701141313H1	SOYMON038	g974781	BLASTN	827	1e-60	83
2894	550	701012547H1	SOYMON019	g1814402	BLASTN	831	1e-60	83
2895	550	701008558H1	SOYMON019	g1814402	BLASTN	831	1e-60	83
2896	550	700902022H1	SOYMON027	g2738247	BLASTN	831	1e-60	82
2897	550	700959515H1	SOYMON022	g886470	BLASTN	832	1e-60	81
2898	550	701042630H1	SOYMON029	g1814402	BLASTN	819	1e-59	81
2899	550	700941292H1	SOYMON024	g2738247	BLASTN	820	1e-59	81
2900	550	700788526H1	SOYMON011	g974781	BLASTN	491	1e-58	82
2901	550	700894839H1	SOYMON024	g1814402	BLASTN	498	1e-58	82
2902	550	700865873H1	SOYMON016	g974781	BLASTN	808	1e-58	85
2903	550	701015435H1	SOYMON019	g886470	BLASTN	809	1e-58	80
2904	550	700755960H1	SOYMON014	g2738247	BLASTN	809	1e-58	78
2905	550	700876051H1	SOYMON018	g886470	BLASTN	434	1e-57	81
2906	550	701041327H1	SOYMON029	g886470	BLASTN	499	1e-57	83
2907	550	701098902H1	SOYMON028	g2738247	BLASTN	767	1e-57	77
2908	550	700853392H1	SOYMON023	g886470	BLASTN	793	1e-57	82
2909	550	700872645H1	SOYMON018	g974781	BLASTN	798	1e-57	83
2910	550	700989675H1	SOYMON011	g2738247	BLASTN	800	1e-57	79
2911	550	700753487H1	SOYMON014	g1814402	BLASTN	589	1e-56	83
2912	550	700736276H1	SOYMON010	g1814402	BLASTN	782	1e-56	79
2913	550	700891361H1	SOYMON024	g1814402	BLASTN	783	1e-56	81
2914	550	700829712H1	SOYMON019	g974781	BLASTN	790	1e-56	80
2915	550	LIB3050-019-Q1-K1-A1	LIB3050	g974781	BLASTN	663	1e-55	81
2916	550	701001013H1	SOYMON018	g1814402	BLASTN	768	1e-55	84
2917	550	LIB3028-031-Q1-B1-G12	LIB3028	g886470	BLASTN	775	1e-55	82
2918	550	701212782H1	SOYMON035	g886470	BLASTN	621	1e-54	82
2919	550	701008695H1	SOYMON019	g974781	BLASTN	718	1e-54	80
2920	550	700990972H1	SOYMON011	g1814402	BLASTN	766	1e-54	81
2921	550	700789576H2	SOYMON011	g886470	BLASTN	766	1e-54	80
2922	550	700994266H1	SOYMON011	g1814402	BLASTN	408	1e-53	85
2923	550	700731985H1	SOYMON010	g974781	BLASTN	590	1e-53	81
2924	550	700907927H1	SOYMON022	g886470	BLASTN	612	1e-53	80
2925	550	701012079H1	SOYMON019	g974781	BLASTN	669	1e-53	86
2926	550	700753939H1	SOYMON014	g886470	BLASTN	690	1e-53	78
2927	550	701000754H1	SOYMON018	g974781	BLASTN	745	1e-53	76
2928	550	701040287H1	SOYMON029	g886470	BLASTN	747	1e-53	82
2929	550	700891329H1	SOYMON024	g974781	BLASTN	749	1e-53	79
2930	550	700897258H1	SOYMON027	g886470	BLASTN	752	1e-53	86
2931	550	700944949H1	SOYMON024	g974781	BLASTN	426	1e-52	82
2932	550	701108671H1	SOYMON036	g1814402	BLASTN	731	1e-52	77
2933	550	700905233H1	SOYMON022	g886470	BLASTN	732	1e-52	77
2934	550	700958589H1	SOYMON022	g886470	BLASTN	738	1e-52	80
2935	550	700666436H1	SOYMON005	g2738247	BLASTN	739	1e-52	82
2936	550	700829902H1	SOYMON019	g886470	BLASTN	740	1e-52	82
2937	550	700740110H1	SOYMON012	g974781	BLASTN	742	1e-52	83
2938	550	700989055H1	SOYMON011	g886470	BLASTN	538	1e-50	79
2939	550	701213370H1	SOYMON035	g974781	BLASTN	599	1e-50	83
2940	550	701098072H1	SOYMON028	g974781	BLASTN	709	1e-50	74
2941	550	700896128H1	SOYMON027	g886470	BLASTN	714	1e-50	83
2942	550	701060755H1	SOYMON033	g974781	BLASTN	716	1e-50	74
2943	550	700953594H1	SOYMON022	g1814402	BLASTN	695	1e-49	78
2944	550	701046911H1	SOYMON032	g2738247	BLASTN	687	1e-48	81

2945	550	701065707H1	SOYMON034	g1814402	BLASTN	443	1e-47	84
2946	550	700962114H1	SOYMON022	g886470	BLASTN	678	1e-47	84
2947	550	700831826H1	SOYMON019	g2738247	BLASTN	682	1e-47	77
2948	550	701054296H1	SOYMON032	g886470	BLASTN	454	1e-46	83
2949	550	700888738H1	SOYMON024	g974781	BLASTN	552	1e-46	77
2950	550	700892022H1	SOYMON024	g886470	BLASTN	605	1e-46	81
2951	550	700890275H1	SOYMON024	g2738247	BLASTN	666	1e-46	77
2952	550	LIB3051-006-Q1-K1-G9	LIB3051	g974781	BLASTN	670	1e-45	80
2953	550	700889113H1	SOYMON024	g886470	BLASTN	582	1e-43	81
2954	550	700952720H1	SOYMON022	g886470	BLASTN	611	1e-42	76
2955	550	701014761H1	SOYMON019	g886470	BLASTN	318	1e-41	84
2956	550	700753882H1	SOYMON014	g886470	BLASTN	381	1e-41	79
2957	550	700743792H1	SOYMON012	g1814402	BLASTN	610	1e-41	86
2958	550	700990963H1	SOYMON011	g886470	BLASTN	578	1e-39	77
2959	550	700941880H1	SOYMON024	g2738247	BLASTN	583	1e-39	81
2960	550	700898962H1	SOYMON027	g2738247	BLASTN	571	1e-38	80
2961	550	700990865H1	SOYMON011	g2738247	BLASTN	467	1e-37	77
2962	550	700565779H1	SOYMON002	g886470	BLASTN	557	1e-37	70
2963	550	700993903H1	SOYMON011	g974781	BLASTN	562	1e-37	84
2964	550	700941589H1	SOYMON024	g2738247	BLASTN	550	1e-36	81
2965	550	701052554H1	SOYMON032	g886470	BLASTN	534	1e-35	67
2966	550	700991055H1	SOYMON011	g886470	BLASTN	356	1e-34	79
2967	550	701010438H1	SOYMON019	g2738247	BLASTN	508	1e-33	79
2968	550	700756634H1	SOYMON014	g2738247	BLASTN	494	1e-32	76
2969	550	701042980H1	SOYMON029	g886470	BLASTN	461	1e-29	83
2970	550	700682940H1	SOYMON008	g2738247	BLASTN	466	1e-29	83
2971	550	701049575H1	SOYMON032	g886470	BLASTN	433	1e-27	82
2972	550	700982552H1	SOYMON009	g886470	BLASTN	356	1e-25	78
2973	550	700675637H1	SOYMON007	g886470	BLASTN	375	1e-25	79
2974	550	701142153H1	SOYMON038	g886470	BLASTN	377	1e-22	76
2975	550	700682724H1	SOYMON008	g974781	BLASTN	361	1e-19	88
2976	550	701051764H1	SOYMON032	g1814402	BLASTN	211	1e-17	80
2977	550	700867241H1	SOYMON016	g2738248	BLASTX	152	1e-13	88
2978	550	701054954H1	SOYMON032	g2738248	BLASTX	138	1e-12	86
2979	550	700790450H2	SOYMON011	g974781	BLASTN	238	1e-10	80
2980	550	700653979H1	SOYMON003	g1814403	BLASTX	118	1e-9	92
2981	550	700894218H1	SOYMON024	g2738248	BLASTX	122	1e-9	78
2982	550	700863078H1	SOYMON022	g2738247	BLASTN	236	1e-8	78
2983	5758	701209304H1	SOYMON035	g886470	BLASTN	766	1e-54	84
2984	5758	701106455H1	SOYMON036	g886470	BLASTN	723	1e-51	82
2985	5758	700833538H1	SOYMON019	g1814402	BLASTN	609	1e-43	83
2986	5758	701051425H1	SOYMON032	g886470	BLASTN	629	1e-43	83
2987	5758	700654506H1	SOYMON004	g886470	BLASTN	438	1e-26	70
2988	5758	701047795H1	SOYMON032	g974782	BLASTX	161	1e-15	100
2989	5758	701202409H1	SOYMON035	g1814402	BLASTN	310	1e-15	79
2990	8266	700558628H1	SOYMON001	g886470	BLASTN	780	1e-56	74
2991	8266	701207720H1	SOYMON035	g1814402	BLASTN	766	1e-54	74
2992	8266	700557429H1	SOYMON001	g1814402	BLASTN	728	1e-51	75

ADENOSYLHOMOCYSTEINASE (EC 3.3.1.1)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1354	-700154280	700154280H1	SATMON007	g170772	BLASTN	229	1e-23	77

1355	-L30594291	LIB3059-032-Q1-K1-A8	LIB3059	g170772	BLASTN	516	1e-74	73
1356	-L30664307	LIB3066-047-Q1-K1-E7	LIB3066	g170772	BLASTN	628	1e-49	70
1357	503	LIB3079-013-Q1-K1-D3	LIB3079	g170772	BLASTN	1505	1e-139	89
1358	503	LIB3062-017-Q1-K1-A10	LIB3062	g170772	BLASTN	1571	1e-129	89
1359	503	LIB3067-001-Q1-K1-D11	LIB3067	g170772	BLASTN	1654	1e-129	89
1360	503	LIB148-060-Q1-E1-B9	LIB148	g170772	BLASTN	1620	1e-126	90
1361	503	LIB3069-043-Q1-K1-G4	LIB3069	g170772	BLASTN	1360	1e-124	84
1362	503	LIB143-006-Q1-E1-F3	LIB143	g170772	BLASTN	831	1e-120	88
1363	503	LIB189-009-Q1-E1-E9	LIB189	g170772	BLASTN	1551	1e-120	90
1364	503	LIB3060-003-Q1-K1-F9	LIB3060	g170772	BLASTN	1538	1e-119	91
1365	503	700089023H1	SATMON011	g170772	BLASTN	1495	1e-115	91
1366	503	LIB143-061-Q1-E1-C9	LIB143	g170772	BLASTN	1474	1e-114	90
1367	503	LIB3069-034-Q1-K1-E5	LIB3069	g170772	BLASTN	1483	1e-114	88
1368	503	LIB143-061-Q1-E1-E5	LIB143	g170772	BLASTN	1440	1e-113	89
1369	503	LIB3067-040-Q1-K1-H5	LIB3067	g170772	BLASTN	1467	1e-113	88
1370	503	LIB3067-048-Q1-K1-A11	LIB3067	g170772	BLASTN	1207	1e-109	86
1371	503	LIB3068-050-Q1-K1-G6	LIB3068	g170772	BLASTN	1279	1e-109	86
1372	503	LIB3069-036-Q1-K1-F6	LIB3069	g170772	BLASTN	1310	1e-109	92
1373	503	LIB3066-047-Q1-K1-H5	LIB3066	g170772	BLASTN	1353	1e-108	86
1374	503	LIB3059-011-Q1-K1-A5	LIB3059	g170772	BLASTN	1401	1e-107	90
1375	503	LIB3067-056-Q1-K1-E11	LIB3067	g170772	BLASTN	957	1e-106	86
1376	503	LIB189-010-Q1-E1-E12	LIB189	g170772	BLASTN	1144	1e-106	88
1377	503	700084426H1	SATMON011	g170772	BLASTN	1368	1e-105	91
1378	503	700086273H1	SATMON011	g170772	BLASTN	1364	1e-104	91
1379	503	700573027H1	SATMON030	g170772	BLASTN	939	1e-103	89
1380	503	700209360H1	SATMON016	g170772	BLASTN	1350	1e-103	89
1381	503	700619916H1	SATMON034	g170772	BLASTN	1050	1e-102	89
1382	503	700086051H1	SATMON011	g170772	BLASTN	1336	1e-102	90
1383	503	LIB3069-034-Q1-K1-C8	LIB3069	g170772	BLASTN	1322	1e-101	86
1384	503	700026324H1	SATMON003	g170772	BLASTN	1328	1e-101	91
1385	503	700104549H1	SATMON010	g170772	BLASTN	836	1e-100	88
1386	503	700622108H1	SATMON034	g170772	BLASTN	1158	1e-100	88
1387	503	700093980H1	SATMON008	g170772	BLASTN	1312	1e-100	90
1388	503	700077427H1	SATMON007	g170772	BLASTN	1317	1e-100	90
1389	503	700095389H1	SATMON008	g170772	BLASTN	1302	1e-99	91
1390	503	LIB3067-055-Q1-K1-D3	LIB3067	g170772	BLASTN	762	1e-98	87
1391	503	LIB3060-054-Q1-K1-F6	LIB3060	g170772	BLASTN	1009	1e-98	83
1392	503	700083339H1	SATMON011	g170772	BLASTN	1282	1e-98	91
1393	503	700102631H1	SATMON010	g170772	BLASTN	1283	1e-98	89
1394	503	700265625H1	SATMON017	g170772	BLASTN	1286	1e-98	90
1395	503	700095002H1	SATMON008	g170772	BLASTN	1289	1e-98	88
1396	503	700094761H1	SATMON008	g170772	BLASTN	1289	1e-98	88
1397	503	700073832H1	SATMON007	g170772	BLASTN	1270	1e-97	88
1398	503	700047817H1	SATMON003	g170772	BLASTN	1272	1e-97	91
1399	503	700091149H1	SATMON011	g170772	BLASTN	1262	1e-96	89
1400	503	700098584H1	SATMON009	g170772	BLASTN	1264	1e-96	91
1401	503	700085932H1	SATMON011	g170772	BLASTN	1266	1e-96	89
1402	503	700094081H1	SATMON008	g170772	BLASTN	1269	1e-96	91
1403	503	700202442H1	SATMON003	g170772	BLASTN	1145	1e-95	88
1404	503	700049770H1	SATMON003	g170772	BLASTN	1246	1e-95	89
1405	503	LIB3079-020-Q1-K1-B12	LIB3079	g170772	BLASTN	1249	1e-95	85
1406	503	700090226H1	SATMON011	g170772	BLASTN	1252	1e-95	90
1407	503	700088191H1	SATMON011	g170772	BLASTN	1253	1e-95	90
1408	503	700076743H1	SATMON007	g170772	BLASTN	1256	1e-95	90

1409	503	LIB189-009-Q1-E1-E10	LIB189	g170772	BLASTN	789	1e-94	88
1410	503	700072038H1	SATMON007	g170772	BLASTN	1237	1e-94	89
1411	503	700082967H1	SATMON011	g170772	BLASTN	1239	1e-94	89
1412	503	LIB3068-062-Q1-K1-A3	LIB3068	g170772	BLASTN	1148	1e-93	82
1413	503	700094713H1	SATMON008	g170772	BLASTN	1222	1e-93	88
1414	503	700095620H1	SATMON008	g170772	BLASTN	1173	1e-92	89
1415	503	700242509H1	SATMON010	g170772	BLASTN	1213	1e-92	92
1416	503	700071923H1	SATMON007	g170772	BLASTN	1214	1e-92	90
1417	503	700575314H1	SATMON030	g170772	BLASTN	1149	1e-91	88
1418	503	700086654H1	SATMON011	g170772	BLASTN	1199	1e-91	90
1419	503	700241072H1	SATMON010	g170772	BLASTN	1205	1e-91	91
1420	503	700047361H1	SATMON003	g170772	BLASTN	1110	1e-90	89
1421	503	700217056H1	SATMON016	g170772	BLASTN	1187	1e-90	91
1422	503	LIB3067-005-Q1-K1-F2	LIB3067	g170772	BLASTN	1188	1e-90	84
1423	503	700075495H1	SATMON007	g170772	BLASTN	1193	1e-90	86
1424	503	700084783H1	SATMON011	g170772	BLASTN	1194	1e-90	91
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1426	503	700448818H1	SATMON028	g170772	BLASTN	1174	1e-89	91
1427	503	700159079H1	SATMON012	g170772	BLASTN	1177	1e-89	92
1428	503	700094408H1	SATMON008	g170772	BLASTN	1177	1e-89	92
1429	503	700348440H1	SATMON023	g170772	BLASTN	1182	1e-89	89
1430	503	700077082H1	SATMON007	g170772	BLASTN	1182	1e-89	92
1431	503	700082111H1	SATMON011	g170772	BLASTN	1183	1e-89	89
1432	503	700239752H1	SATMON010	g170772	BLASTN	1164	1e-88	89
1433	503	700029456H1	SATMON003	g170772	BLASTN	1166	1e-88	90
1434	503	700209430H1	SATMON016	g170772	BLASTN	1170	1e-88	90
1435	503	700208660H1	SATMON016	g170772	BLASTN	930	1e-87	90
1436	503	700213138H1	SATMON016	g170772	BLASTN	1042	1e-87	89
1437	503	700102234H1	SATMON010	g170772	BLASTN	634	1e-85	90
1438	503	700450479H1	SATMON028	g170772	BLASTN	1015	1e-85	90
1439	503	700095372H1	SATMON008	g170772	BLASTN	1126	1e-85	90
1440	503	700218734H1	SATMON011	g170772	BLASTN	1128	1e-85	94
1441	503	700256858H1	SATMON017	g170772	BLASTN	1133	1e-85	90
1442	503	700221063H1	SATMON011	g170772	BLASTN	1137	1e-85	90
1443	503	700105439H1	SATMON010	g170772	BLASTN	1118	1e-84	89
1444	503	700085174H1	SATMON011	g170772	BLASTN	1125	1e-84	90
1445	503	700242421H1	SATMON010	g170772	BLASTN	1125	1e-84	92
1446	503	700077492H1	SATMON007	g170772	BLASTN	774	1e-83	88
1447	503	700209295H1	SATMON016	g170772	BLASTN	914	1e-83	89
1448	503	700455826H1	SATMON029	g170772	BLASTN	978	1e-83	91
1449	503	700213370H1	SATMON016	g170772	BLASTN	1110	1e-83	92
1450	503	700352095H1	SATMON023	g170772	BLASTN	1111	1e-83	89
1451	503	700076842H1	SATMON007	g170772	BLASTN	1112	1e-83	90
1452	503	700215872H1	SATMON016	g170772	BLASTN	1113	1e-83	89
1453	503	700073645H1	SATMON007	g170772	BLASTN	1113	1e-83	89
1454	503	700048153H1	SATMON003	g170772	BLASTN	651	1e-82	90
1455	503	700073307H1	SATMON007	g170772	BLASTN	762	1e-82	90
1456	503	700350009H1	SATMON023	g170772	BLASTN	924	1e-82	88
1457	503	LIB143-059-Q1-E1-C5	LIB143	g170772	BLASTN	1029	1e-82	83
1458	503	700073955H1	SATMON007	g170772	BLASTN	1090	1e-82	90
1459	503	LIB3059-042-Q1-K1-H12	LIB3059	g170772	BLASTN	1090	1e-82	85
1460	503	700238024H1	SATMON010	g170772	BLASTN	1091	1e-82	90
1461	503	700155863H1	SATMON007	g170772	BLASTN	1091	1e-82	93
1462	503	700217890H1	SATMON016	g170772	BLASTN	1093	1e-82	91

1463	503	700239314H1	SATMON010	g170772	BLASTN	1094	1e-82	88
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1465	503	700235469H1	SATMON010	g170772	BLASTN	1098	1e-82	91
1466	503	700164224H1	SATMON013	g170772	BLASTN	710	1e-81	92
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1469	503	700159326H1	SATMON012	g170772	BLASTN	777	1e-80	91
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1473	503	700242676H1	SATMON010	g170772	BLASTN	940	1e-80	89
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1477	503	700072491H1	SATMON007	g170772	BLASTN	604	1e-79	88
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1479	503	700451515H1	SATMON028	g170772	BLASTN	1057	1e-79	85
1480	503	700468929H1	SATMON025	g170772	BLASTN	1061	1e-79	84
1481	503	700205801H1	SATMON003	g170772	BLASTN	1064	1e-79	91
1482	503	700611326H1	SATMON022	g170772	BLASTN	725	1e-78	88
1483	503	700082291H1	SATMON011	g170772	BLASTN	1043	1e-78	87
1484	503	700071695H1	SATMON007	g170772	BLASTN	1048	1e-78	89
1485	503	700551561H1	SATMON022	g170772	BLASTN	746	1e-77	88
1486	503	700221019H1	SATMON011	g170772	BLASTN	856	1e-77	89
1487	503	LIB3078-007-Q1-K1-C5	LIB3078	g170772	BLASTN	936	1e-77	83
1488	503	700049925H1	SATMON003	g170772	BLASTN	954	1e-77	89
1489	503	700154101H1	SATMON007	g170772	BLASTN	1036	1e-77	88
1490	503	700380151H1	SATMON021	g170772	BLASTN	613	1e-76	88
1491	503	700243831H1	SATMON010	g170772	BLASTN	842	1e-76	89
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1493	503	700087847H1	SATMON011	g170772	BLASTN	1009	1e-75	82
1494	503	700208747H1	SATMON016	g170772	BLASTN	1010	1e-75	89
1495	503	700071795H1	SATMON007	g170772	BLASTN	1011	1e-75	89
1496	503	700157002H1	SATMON012	g170772	BLASTN	1012	1e-75	90
1497	503	700201608H1	SATMON003	g170772	BLASTN	1015	1e-75	85
1498	503	700451541H1	SATMON028	g170772	BLASTN	1015	1e-75	86
1499	503	700212182H1	SATMON016	g170772	BLASTN	1015	1e-75	88
1500	503	700381485H1	SATMON023	g170772	BLASTN	1017	1e-75	91
1501	503	700096793H1	SATMON008	g170772	BLASTN	761	1e-74	87
1502	503	700093025H1	SATMON008	g170772	BLASTN	996	1e-74	89
1503	503	700017129H1	SATMON001	g170772	BLASTN	998	1e-74	92
1504	503	700216525H1	SATMON016	g170772	BLASTN	1000	1e-74	86
1505	503	700087912H1	SATMON011	g170772	BLASTN	1001	1e-74	91
1506	503	700172618H1	SATMON013	g170772	BLASTN	1001	1e-74	91
1507	503	700801894H1	SATMON036	g170772	BLASTN	1002	1e-74	90
1508	503	700162040H1	SATMON012	g170772	BLASTN	1003	1e-74	92
1509	503	700212084H1	SATMON016	g170772	BLASTN	1004	1e-74	89
1510	503	700027971H1	SATMON003	g170772	BLASTN	982	1e-73	92
1511	503	700077321H1	SATMON007	g170772	BLASTN	985	1e-73	89
1512	503	700619884H1	SATMON034	g170772	BLASTN	987	1e-73	87
1513	503	700457201H1	SATMON029	g170772	BLASTN	988	1e-73	87
1514	503	700082970H1	SATMON011	g170772	BLASTN	990	1e-73	89
1515	503	700083350H1	SATMON011	g170772	BLASTN	971	1e-72	89
1516	503	700017732H1	SATMON001	g170772	BLASTN	973	1e-72	88

1517	503	700094688H1	SATMON008	g170772	BLASTN	975	1e-72	89
1518	503	700170985H1	SATMON013	g170772	BLASTN	979	1e-72	89
1519	503	700451847H1	SATMON028	g170772	BLASTN	872	1e-71	84
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1521	503	700106189H1	SATMON010	g170772	BLASTN	960	1e-71	88
1522	503	700160075H1	SATMON012	g170772	BLASTN	968	1e-71	90
1523	503	700084952H1	SATMON011	g170772	BLASTN	552	1e-70	78
1524	503	700348238H1	SATMON023	g170772	BLASTN	946	1e-70	87
1525	503	700151310H1	SATMON007	g170772	BLASTN	948	1e-70	90
1526	503	700048962H1	SATMON003	g170772	BLASTN	677	1e-69	88
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1530	503	700073939H1	SATMON007	g170772	BLASTN	931	1e-68	88
1531	503	700027715H1	SATMON003	g170772	BLASTN	472	1e-67	90
1532	503	700072843H1	SATMON007	g170772	BLASTN	692	1e-67	87
1533	503	700439356H1	SATMON026	g170772	BLASTN	777	1e-67	85
1534	503	700073556H1	SATMON007	g170772	BLASTN	910	1e-67	88
1535	503	700154461H1	SATMON007	g170772	BLASTN	911	1e-67	87
1536	503	700201909H1	SATMON003	g170772	BLASTN	912	1e-67	88
1537	503	700104384H1	SATMON010	g170772	BLASTN	665	1e-66	89
1538	503	700152707H1	SATMON007	g170772	BLASTN	898	1e-66	88
1539	503	700076625H1	SATMON007	g170772	BLASTN	900	1e-66	85
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1541	503	700457294H1	SATMON029	g170772	BLASTN	446	1e-65	88
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1543	503	700072428H2	SATMON007	g170772	BLASTN	890	1e-65	88
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1545	503	700072443H2	SATMON007	g170772	BLASTN	894	1e-65	89
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1547	503	700240615H1	SATMON010	g170772	BLASTN	877	1e-64	88
1548	503	700155233H1	SATMON007	g170772	BLASTN	883	1e-64	90
1549	503	700241210H1	SATMON010	g170772	BLASTN	822	1e-63	87
1550	503	700478070H1	SATMON025	g170772	BLASTN	863	1e-63	82
1551	503	700447433H1	SATMON027	g170772	BLASTN	865	1e-63	84
1552	503	700215884H1	SATMON016	g170772	BLASTN	865	1e-63	88
1553	503	700575250H1	SATMON030	g170772	BLASTN	820	1e-61	84
1554	503	700209283H1	SATMON016	g170772	BLASTN	833	1e-60	86
1555	503	700213479H1	SATMON016	g170772	BLASTN	834	1e-60	88
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1558	503	700151036H1	SATMON007	g170772	BLASTN	816	1e-59	91
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1561	503	700618168H1	SATMON033	g170772	BLASTN	385	1e-58	80
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1563	503	700224131H1	SATMON011	g170772	BLASTN	794	1e-57	88
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1565	503	700027716H1	SATMON003	g170772	BLASTN	642	1e-56	79
1566	503	700073305H1	SATMON007	g170772	BLASTN	787	1e-56	83
1567	503	700257376H1	SATMON017	g170772	BLASTN	362	1e-55	87
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1569	503	700219086H1	SATMON011	g170772	BLASTN	769	1e-55	87
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1571	503	700424527H1	SATMONN01	g170772	BLASTN	777	1e-55	83
1572	503	700073183H1	SATMON007	g170772	BLASTN	607	1e-54	85
1573	503	700217412H1	SATMON016	g170772	BLASTN	755	1e-54	87
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1576	503	700446645H1	SATMON027	g170772	BLASTN	744	1e-53	87
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1578	503	700617325H1	SATMON033	g170772	BLASTN	751	1e-53	91
1579	503	700239901H1	SATMON010	g170772	BLASTN	641	1e-52	86
1580	503	700155130H1	SATMON007	g170772	BLASTN	646	1e-52	88
1581	503	700083360H1	SATMON011	g170772	BLASTN	739	1e-52	87
1582	503	700236915H1	SATMON010	g170772	BLASTN	740	1e-52	87
1583	503	700074780H1	SATMON007	g170772	BLASTN	528	1e-51	83
1584	503	700479515H1	SATMON034	g170772	BLASTN	718	1e-51	88
1585	503	700215545H1	SATMON016	g170772	BLASTN	724	1e-51	90
1586	503	700165354H1	SATMON013	g170772	BLASTN	728	1e-51	86
1587	503	700353994H1	SATMON024	g170772	BLASTN	713	1e-50	87
1588	503	700153619H1	SATMON007	g170772	BLASTN	694	1e-49	86
1589	503	700074779H1	SATMON007	g170772	BLASTN	705	1e-49	77
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1591	503	700221207H1	SATMON011	g170772	BLASTN	692	1e-48	86
1592	503	700264257H1	SATMON017	g170772	BLASTN	672	1e-47	87
1593	503	700152216H1	SATMON007	g170772	BLASTN	659	1e-46	88
1594	503	700150643H1	SATMON007	g170772	BLASTN	659	1e-46	88
1595	503	700156027H1	SATMON007	g170772	BLASTN	655	1e-45	93
1596	503	700260335H1	SATMON017	g170772	BLASTN	385	1e-44	79
1597	503	700623795H1	SATMON034	g170772	BLASTN	401	1e-44	88
1598	503	700150581H1	SATMON007	g170772	BLASTN	640	1e-44	88
1599	503	700151970H1	SATMON007	g170772	BLASTN	645	1e-44	87
1600	503	700151780H1	SATMON007	g170772	BLASTN	628	1e-43	88
1601	503	700575547H1	SATMON030	g170772	BLASTN	614	1e-42	90
1602	503	LIB143-026-Q1-E1-A5	LIB143	g170772	BLASTN	633	1e-42	77
1603	503	700347945H1	SATMON023	g170772	BLASTN	586	1e-40	84
1604	503	700074416H1	SATMON007	g170772	BLASTN	589	1e-40	86
1605	503	700352990H1	SATMON024	g170772	BLASTN	575	1e-39	92
1606	503	700156025H1	SATMON007	g170772	BLASTN	568	1e-38	91
1607	503	700432072H1	SATMONN01	g170772	BLASTN	374	1e-37	84
1608	503	700354249H1	SATMON024	g170772	BLASTN	528	1e-35	93
1609	503	700617977H1	SATMON033	g170772	BLASTN	535	1e-35	87
1610	503	700218312H1	SATMON016	g170772	BLASTN	236	1e-33	91
1611	503	700456080H1	SATMON029	g170772	BLASTN	513	1e-33	83
1612	503	700349395H1	SATMON023	g170772	BLASTN	500	1e-32	91
1613	503	700051637H1	SATMON003	g170772	BLASTN	249	1e-31	86
1614	503	700202153H1	SATMON003	g170772	BLASTN	474	1e-30	91
1615	503	700256951H1	SATMON017	g170772	BLASTN	454	1e-29	86
1616	503	700150542H1	SATMON007	g170772	BLASTN	343	1e-28	76
1617	503	700151629H1	SATMON007	g170772	BLASTN	445	1e-28	85
1618	503	700446654H1	SATMON027	g170772	BLASTN	437	1e-27	84
1619	503	700155814H1	SATMON007	g170772	BLASTN	428	1e-26	88
1620	503	700161019H1	SATMON012	g2588780	BLASTN	417	1e-25	92
1621	503	700377236H1	SATMON019	g170772	BLASTN	402	1e-24	84
1622	503	LIB3067-036-Q1-K1-A4	LIB3067	g1220121	BLASTN	224	1e-22	84
1623	503	700158933H1	SATMON012	g170772	BLASTN	368	1e-21	90
1624	503	700155365H1	SATMON007	g170772	BLASTN	347	1e-20	91

1625	503	700405366H1	SATMON029	g170772	BLASTN	311	1e-17	88
1626	503	700159812H1	SATMON012	g407412	BLASTX	160	1e-15	96
1627	503	700209759H1	SATMON016	g170772	BLASTN	239	1e-15	90
1628	503	700154527H1	SATMON007	g170772	BLASTN	250	1e-12	92
1629	503	700449637H1	SATMON028	g170772	BLASTN	201	1e-10	78
1630	503	700096829H1	SATMON008	g170772	BLASTN	216	1e-9	89
2993	-700661285	700661285H1	SOYMON005	g1857024	BLASTX	95	1e-12	100
2994	-700750570	700750570H1	SOYMON014	g170772	BLASTN	414	1e-24	81
2995	-700752735	700752735H1	SOYMON014	g170772	BLASTN	446	1e-27	78
2996	-700755052	700755052H1	SOYMON014	g170772	BLASTN	547	1e-45	74
2997	-700756501	700756501H1	SOYMON014	g535583	BLASTN	717	1e-50	84
2998	-700831127	700831127H1	SOYMON019	g535583	BLASTN	862	1e-63	83
2999	-700851779	700851779H1	SOYMON023	g170772	BLASTN	505	1e-33	77
3000	-700888715	700888715H1	SOYMON024	g535583	BLASTN	442	1e-31	91
3001	-700889420	700889420H1	SOYMON024	g1220121	BLASTN	893	1e-65	84
3002	-700895218	700895218H1	SOYMON024	g407411	BLASTN	816	1e-59	83
3003	-700941379	700941379H1	SOYMON024	g170772	BLASTN	424	1e-33	72
3004	-700986855	700986855H1	SOYMON009	g170772	BLASTN	701	1e-49	74
3005	-701070484	701070484H1	SOYMON034	g407411	BLASTN	362	1e-43	73
3006	-701136279	701136279H1	SOYMON038	g170772	BLASTN	651	1e-56	80
3007	-GM16478	LIB3054-007-Q1-N1-G3	LIB3054	g2244750	BLASTX	70	1e-27	61
3008	-GM23819	LIB3040-019-Q1-E1-C5	LIB3040	g535583	BLASTN	258	1e-10	88
3009	-GM29758	LIB3050-016-Q1-E1-B11	LIB3050	g535583	BLASTN	391	1e-42	70
3010	16	LIB3030-003-Q1-B1-B11	LIB3030	g3088578	BLASTN	1577	1e-122	87
3011	16	LIB3050-023-Q1-K1-H9	LIB3050	g535583	BLASTN	1498	1e-116	86
3012	16	LIB3030-003-Q1-B1-F7	LIB3030	g535583	BLASTN	1469	1e-113	86
3013	16	LIB3055-005-Q1-N1-C11	LIB3055	g170772	BLASTN	1205	1e-109	84
3014	16	LIB3065-011-Q1-N1-A3	LIB3065	g170772	BLASTN	622	1e-107	88
3015	16	700652256H1	SOYMON003	g170772	BLASTN	591	1e-90	88
3016	16	LIB3065-011-Q1-N1-A4	LIB3065	g170772	BLASTN	1182	1e-89	78
3017	16	700653827H1	SOYMON003	g170772	BLASTN	660	1e-87	81
3018	16	701099940H1	SOYMON028	g1220121	BLASTN	1154	1e-87	88
3019	16	701003671H1	SOYMON019	g170772	BLASTN	1131	1e-85	92
3020	16	700752105H1	SOYMON014	g170772	BLASTN	1116	1e-84	91
3021	16	700653057H1	SOYMON003	g170772	BLASTN	990	1e-83	88
3022	16	700945531H1	SOYMON024	g535583	BLASTN	1109	1e-83	89
3023	16	700980013H1	SOYMON009	g535583	BLASTN	1109	1e-83	88
3024	16	701127812H1	SOYMON037	g170772	BLASTN	1110	1e-83	91
3025	16	LIB3056-014-Q1-N1-F8	LIB3056	g170772	BLASTN	798	1e-81	82
3026	16	700653862H1	SOYMON003	g1220121	BLASTN	960	1e-80	88
3027	16	700994148H1	SOYMON011	g535583	BLASTN	1069	1e-80	86
3028	16	700984184H1	SOYMON009	g170772	BLASTN	756	1e-79	86
3029	16	701123715H1	SOYMON037	g1220121	BLASTN	1059	1e-79	87
3030	16	700839038H1	SOYMON020	g170772	BLASTN	1060	1e-79	93
3031	16	700978445H1	SOYMON009	g170772	BLASTN	1062	1e-79	89
3032	16	701123035H1	SOYMON037	g170772	BLASTN	829	1e-78	90
3033	16	701041545H1	SOYMON029	g170772	BLASTN	1030	1e-77	86
3034	16	700898192H1	SOYMON027	g1220121	BLASTN	1031	1e-77	89
3035	16	700985750H1	SOYMON009	g169662	BLASTN	1033	1e-77	85
3036	16	700730995H1	SOYMON009	g170772	BLASTN	1040	1e-77	91
3037	16	700555909H1	SOYMON001	g535583	BLASTN	593	1e-76	84
3038	16	700746538H1	SOYMON013	g535583	BLASTN	834	1e-76	88
3039	16	700941380H1	SOYMON024	g170772	BLASTN	991	1e-76	91
3040	16	701118851H1	SOYMON037	g170772	BLASTN	1019	1e-76	89

3041	16	701065379H1	SOYMON034	g535583	BLASTN	1022	1e-76	86
3042	16	701209645H1	SOYMON035	g170772	BLASTN	795	1e-75	86
3043	16	701055017H1	SOYMON032	g1220121	BLASTN	877	1e-75	86
3044	16	701015213H1	SOYMON019	g535583	BLASTN	1008	1e-75	87
3045	16	700982770H1	SOYMON009	g1220121	BLASTN	1009	1e-75	85
3046	16	701212420H1	SOYMON035	g535583	BLASTN	1012	1e-75	86
3047	16	700977916H1	SOYMON009	g170772	BLASTN	700	1e-74	89
3048	16	700974401H1	SOYMON005	g1220121	BLASTN	857	1e-74	89
3049	16	700645749H1	SOYMON010	g170772	BLASTN	996	1e-74	82
3050	16	700646620H1	SOYMON014	g170772	BLASTN	996	1e-74	89
3051	16	701126103H1	SOYMON037	g170772	BLASTN	1000	1e-74	88
3052	16	700978001H1	SOYMON009	g170772	BLASTN	563	1e-73	89
3053	16	700561987H1	SOYMON002	g1220121	BLASTN	782	1e-73	87
3054	16	701101526H1	SOYMON028	g1220121	BLASTN	810	1e-73	88
3055	16	700562680H1	SOYMON002	g170772	BLASTN	985	1e-73	87
3056	16	700560521H1	SOYMON001	g170772	BLASTN	988	1e-73	80
3057	16	701055544H1	SOYMON032	g170772	BLASTN	992	1e-73	88
3058	16	701061418H1	SOYMON033	g170772	BLASTN	993	1e-73	87
3059	16	701049514H1	SOYMON032	g1220121	BLASTN	884	1e-72	88
3060	16	700646215H1	SOYMON012	g170772	BLASTN	971	1e-72	89
3061	16	701014875H1	SOYMON019	g535583	BLASTN	973	1e-72	87
3062	16	700874718H1	SOYMON018	g169662	BLASTN	974	1e-72	86
3063	16	700897796H1	SOYMON027	g1220121	BLASTN	977	1e-72	87
3064	16	700548019H1	SOYMON001	g170772	BLASTN	547	1e-71	87
3065	16	700904875H1	SOYMON022	g535583	BLASTN	963	1e-71	88
3066	16	701106315H1	SOYMON036	g170772	BLASTN	557	1e-70	90
3067	16	700745023H1	SOYMON013	g407411	BLASTN	949	1e-70	85
3068	16	701120348H1	SOYMON037	g170772	BLASTN	956	1e-70	86
3069	16	701124508H1	SOYMON037	g170772	BLASTN	578	1e-69	84
3070	16	700956183H1	SOYMON022	g1220121	BLASTN	625	1e-69	87
3071	16	700894619H1	SOYMON024	g535583	BLASTN	828	1e-69	85
3072	16	700892392H1	SOYMON024	g535583	BLASTN	936	1e-69	85
3073	16	700730676H1	SOYMON009	g535583	BLASTN	939	1e-69	86
3074	16	700728913H1	SOYMON009	g407411	BLASTN	940	1e-69	87
3075	16	700845737H1	SOYMON021	g1220121	BLASTN	528	1e-68	88
3076	16	700990961H1	SOYMON011	g170772	BLASTN	582	1e-68	85
3077	16	700983443H1	SOYMON009	g170772	BLASTN	922	1e-68	86
3078	16	701120754H1	SOYMON037	g170772	BLASTN	923	1e-68	88
3079	16	700900486H1	SOYMON027	g1220121	BLASTN	924	1e-68	83
3080	16	701133574H2	SOYMON038	g170772	BLASTN	929	1e-68	88
3081	16	700900924H1	SOYMON027	g170772	BLASTN	931	1e-68	88
3082	16	701056706H1	SOYMON032	g170772	BLASTN	486	1e-67	89
3083	16	701110051H1	SOYMON036	g170772	BLASTN	910	1e-67	88
3084	16	701136325H1	SOYMON038	g170772	BLASTN	915	1e-67	87
3085	16	700750809H1	SOYMON014	g170772	BLASTN	900	1e-66	88
3086	16	700686607H1	SOYMON008	g170772	BLASTN	901	1e-66	88
3087	16	700848261H1	SOYMON021	g535583	BLASTN	905	1e-66	87
3088	16	700686634H1	SOYMON008	g170772	BLASTN	905	1e-66	88
3089	16	700891285H1	SOYMON024	g170772	BLASTN	906	1e-66	88
3090	16	700560291H1	SOYMON001	g170772	BLASTN	906	1e-66	88
3091	16	700752975H1	SOYMON014	g170772	BLASTN	907	1e-66	89
3092	16	701006013H2	SOYMON019	g170772	BLASTN	908	1e-66	87
3093	16	700974038H1	SOYMON005	g1220121	BLASTN	631	1e-65	88
3094	16	701047024H1	SOYMON032	g170772	BLASTN	707	1e-65	89

3095	16	700900409H1	SOYMON027	g170772	BLASTN	737	1e-65	83
3096	16	701137320H1	SOYMON038	g170772	BLASTN	769	1e-65	88
3097	16	700978805H1	SOYMON009	g170772	BLASTN	886	1e-65	84
3098	16	700726195H1	SOYMON009	g1220121	BLASTN	889	1e-65	87
3099	16	700661112H1	SOYMON005	g170772	BLASTN	661	1e-64	85
3100	16	700989712H1	SOYMON011	g170772	BLASTN	788	1e-64	88
3101	16	700752287H1	SOYMON014	g170772	BLASTN	875	1e-64	85
3102	16	700964226H1	SOYMON022	g170772	BLASTN	877	1e-64	88
3103	16	700847346H1	SOYMON021	g170772	BLASTN	880	1e-64	83
3104	16	700756428H1	SOYMON014	g170772	BLASTN	883	1e-64	88
3105	16	701049928H1	SOYMON032	g170772	BLASTN	885	1e-64	82
3106	16	700848652H1	SOYMON021	g170772	BLASTN	477	1e-63	89
3107	16	700898929H1	SOYMON027	g1220121	BLASTN	514	1e-63	87
3108	16	700903523H1	SOYMON022	g170772	BLASTN	527	1e-63	82
3109	16	700983745H1	SOYMON009	g1220121	BLASTN	863	1e-63	88
3110	16	700890587H1	SOYMON024	g170772	BLASTN	865	1e-63	86
3111	16	700969917H1	SOYMON005	g407411	BLASTN	868	1e-63	83
3112	16	700808487H1	SOYMON024	g170772	BLASTN	870	1e-63	88
3113	16	700749968H1	SOYMON013	g170772	BLASTN	871	1e-63	87
3114	16	700751254H1	SOYMON014	g170772	BLASTN	575	1e-62	87
3115	16	701014277H1	SOYMON019	g170772	BLASTN	739	1e-62	86
3116	16	700853635H1	SOYMON023	g170772	BLASTN	854	1e-62	87
3117	16	700752357H1	SOYMON014	g170772	BLASTN	856	1e-62	88
3118	16	700754523H1	SOYMON014	g170772	BLASTN	856	1e-62	92
3119	16	700982153H1	SOYMON009	g170772	BLASTN	400	1e-61	82
3120	16	700958283H1	SOYMON022	g170772	BLASTN	839	1e-61	87
3121	16	700980911H1	SOYMON009	g170772	BLASTN	842	1e-61	82
3122	16	700788112H1	SOYMON011	g170772	BLASTN	844	1e-61	83
3123	16	701005927H1	SOYMON019	g170772	BLASTN	849	1e-61	88
3124	16	700756443H1	SOYMON014	g170772	BLASTN	849	1e-61	88
3125	16	700658914H1	SOYMON004	g1220121	BLASTN	468	1e-60	85
3126	16	701135266H1	SOYMON038	g170772	BLASTN	827	1e-60	87
3127	16	700982179H1	SOYMON009	g170772	BLASTN	827	1e-60	82
3128	16	700754593H1	SOYMON014	g170772	BLASTN	832	1e-60	81
3129	16	700831723H1	SOYMON019	g170772	BLASTN	814	1e-59	91
3130	16	700986775H1	SOYMON009	g170772	BLASTN	815	1e-59	92
3131	16	700755219H1	SOYMON014	g170772	BLASTN	820	1e-59	84
3132	16	701015494H1	SOYMON019	g170772	BLASTN	822	1e-59	88
3133	16	701008473H1	SOYMON019	g170772	BLASTN	823	1e-59	87
3134	16	700754981H1	SOYMON014	g170772	BLASTN	824	1e-59	88
3135	16	700870790H1	SOYMON018	g170772	BLASTN	825	1e-59	81
3136	16	700833069H1	SOYMON019	g170772	BLASTN	806	1e-58	86
3137	16	700680127H2	SOYMON008	g535583	BLASTN	807	1e-58	86
3138	16	701015374H1	SOYMON019	g170772	BLASTN	807	1e-58	87
3139	16	700872895H1	SOYMON018	g535583	BLASTN	808	1e-58	88
3140	16	701137912H1	SOYMON038	g170772	BLASTN	374	1e-57	83
3141	16	700984063H1	SOYMON009	g1220121	BLASTN	575	1e-57	79
3142	16	700991988H1	SOYMON011	g170772	BLASTN	791	1e-57	82
3143	16	700873915H1	SOYMON018	g170772	BLASTN	793	1e-57	88
3144	16	700978721H1	SOYMON009	g170772	BLASTN	797	1e-57	78
3145	16	701213679H1	SOYMON035	g170772	BLASTN	798	1e-57	88
3146	16	701102954H1	SOYMON028	g170772	BLASTN	799	1e-57	79
3147	16	700888289H1	SOYMON024	g170772	BLASTN	712	1e-56	91
3148	16	700962086H1	SOYMON022	g170772	BLASTN	783	1e-56	88

3149	16	701052227H1	SOYMON032	g535583	BLASTN	788	1e-56	86
3150	16	700755177H1	SOYMON014	g170772	BLASTN	768	1e-55	88
3151	16	701123136H1	SOYMON037	g170772	BLASTN	771	1e-55	82
3152	16	700979067H1	SOYMON009	g170772	BLASTN	776	1e-55	90
3153	16	700755513H1	SOYMON014	g170772	BLASTN	759	1e-54	92
3154	16	700756639H1	SOYMON014	g170772	BLASTN	761	1e-54	81
3155	16	700554077H1	SOYMON001	g170772	BLASTN	371	1e-53	86
3156	16	700653194H1	SOYMON003	g170772	BLASTN	395	1e-53	86
3157	16	701110348H1	SOYMON036	g170772	BLASTN	743	1e-53	81
3158	16	700753973H1	SOYMON014	g170772	BLASTN	743	1e-53	87
3159	16	701011009H1	SOYMON019	g535583	BLASTN	733	1e-52	81
3160	16	700739086H1	SOYMON012	g170772	BLASTN	456	1e-51	87
3161	16	700740140H1	SOYMON012	g170772	BLASTN	723	1e-51	89
3162	16	700565040H1	SOYMON002	g170772	BLASTN	729	1e-51	74
3163	16	701148186H1	SOYMON031	g535583	BLASTN	665	1e-50	85
3164	16	701142734H1	SOYMON038	g535583	BLASTN	670	1e-47	86
3165	16	700754441H1	SOYMON014	g170772	BLASTN	385	1e-46	93
3166	16	701102588H1	SOYMON028	g1220121	BLASTN	662	1e-46	89
3167	16	700900656H1	SOYMON027	g535583	BLASTN	635	1e-44	87
3168	16	700974207H1	SOYMON005	g535583	BLASTN	641	1e-44	85
3169	16	700982081H1	SOYMON009	g170772	BLASTN	494	1e-43	78
3170	16	701130026H1	SOYMON037	g1220121	BLASTN	482	1e-42	86
3171	16	701009720H1	SOYMON019	g170772	BLASTN	621	1e-42	87
3172	16	700962602H1	SOYMON022	g170772	BLASTN	357	1e-40	91
3173	16	700724934H1	SOYMON009	g535583	BLASTN	561	1e-40	81
3174	16	700729305H1	SOYMON009	g535583	BLASTN	544	1e-39	82
3175	16	701210054H1	SOYMON035	g535583	BLASTN	569	1e-38	85
3176	16	700790192H1	SOYMON011	g535583	BLASTN	293	1e-37	83
3177	16	700984076H1	SOYMON009	g170772	BLASTN	198	1e-35	88
3178	16	700726562H1	SOYMON009	g170772	BLASTN	307	1e-34	78
3179	16	701211376H1	SOYMON035	g170772	BLASTN	524	1e-34	86
3180	16	700753085H1	SOYMON014	g2588780	BLASTN	358	1e-33	76
3181	16	700727993H1	SOYMON009	g535583	BLASTN	465	1e-32	84
3182	16	700561072H1	SOYMON001	g170772	BLASTN	473	1e-30	83
3183	16	701211464H1	SOYMON035	g170772	BLASTN	464	1e-28	81
3184	16	701098045H1	SOYMON028	g169660	BLASTN	386	1e-21	78
3185	16	700945233H1	SOYMON024	g407412	BLASTX	150	1e-18	87
3186	16	700752655H1	SOYMON014	g758247	BLASTX	172	1e-16	94
3187	16	700735356H1	SOYMON010	g758247	BLASTX	152	1e-14	100
3188	16	700683995H1	SOYMON008	g758247	BLASTX	106	1e-13	90
3189	16	700658760H1	SOYMON004	g1857024	BLASTX	123	1e-13	63
3190	16	700762885H1	SOYMON015	g1857024	BLASTX	134	1e-13	89
3191	16	700755740H1	SOYMON014	g170773	BLASTX	149	1e-13	100
3192	16	700854969H1	SOYMON023	g170772	BLASTN	178	1e-12	80
3193	16	701143036H1	SOYMON038	g169661	BLASTX	113	1e-8	83
3194	18409	700786561H1	SOYMON011	g535583	BLASTN	992	1e-73	85
3195	18409	701008057H1	SOYMON019	g535583	BLASTN	831	1e-60	87
3196	18409	701037442H1	SOYMON029	g535583	BLASTN	669	1e-46	86
3197	18409	700942865H1	SOYMON024	g535583	BLASTN	464	1e-32	86
3198	7322	700651524H1	SOYMON003	g170772	BLASTN	466	1e-65	80
3199	7322	700565758H1	SOYMON002	g170772	BLASTN	450	1e-63	81

CYSTATHIONINE β -SYNTHASE (EC 4.2.1.22)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1631	-700025795	700025795H1	SATMON003	g1323263	BLASTX	186	1e-25	68
1632	20651	700344783H1	SATMON021	g1813975	BLASTX	41	1e-9	53

CYSTATHIONINE γ -LYASE (EC 4.4.1.1)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1633	-700260027	700260027H1	SATMON017	g169475	BLASTX	112	1e-10	75
1634	1228	700027629H1	SATMON003	g169475	BLASTX	189	1e-19	87
3203	-700750583	700750583H1	SOYMON014	g169475	BLASTX	149	1e-13	78
3204	12502	LIB3051-069-Q1-K1-E6	LIB3051	g2641242	BLASTX	86	1e-30	38

O-ACETYLHOMOSERINE (THIOL)-LYASE (EC 4.2.99.10)

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
3200	12502	701135185H1	SOYMON038	g1628606	BLASTX	100	1e-10	48
3201	12502	701042913H1	SOYMON029	g2605905	BLASTX	110	1e-9	42
3202	12502	701059330H1	SOYMON033	g2605905	BLASTX	110	1e-9	42

***Table Headings**

Cluster ID

A cluster ID is arbitrarily assigned to all of those clones which belong to the same cluster at a given stringency and a particular clone will belong to only one cluster at a given stringency. If a cluster contains only a single clone (a “singleton”), then the cluster ID number will be negative, with an absolute value equal to the clone ID number of its single member. The cluster ID entries in the table refer to the cluster with which the particular clone in each row is associated.

Clone ID

The clone ID number refers to the particular clone in the PhytoSeq database. Each clone ID entry in the table refers to the clone whose sequence is used for (1) the sequence comparison whose scores are presented and/or (2) assignment to the particular cluster which is presented. Note that a clone may be included in this table even if its sequence comparison scores fail to meet the minimum standards for similarity. In such a case, the clone is included due solely to its association with a particular cluster for which sequences of one or more other member clones possess the required level of similarity.

Library

The library ID refers to the particular cDNA library from which a given clone is obtained. Each cDNA library is associated with the particular tissue(s), line(s) and developmental stage(s) from which it is isolated.

NCBI gi

Each sequence in the GenBank public database is arbitrarily assigned a unique NCBI gi (National Center for Biotechnology Information GenBank Identifier) number. In this table, the

NCBI gi number which is associated (in the same row) with a given clone refers to the particular GenBank sequence which is used in the sequence comparison. This entry is omitted when a clone is included solely due to its association with a particular cluster.

Method

The entry in the “Method” column of the table refers to the type of BLAST search that is used for the sequence comparison. “CLUSTER” is entered when the sequence comparison scores for a given clone fail to meet the minimum values required for significant similarity. In such cases, the clone is listed in the table solely as a result of its association with a given cluster for which sequences of one or more other member clones possess the required level of similarity.

Score

Each entry in the “Score” column of the table refers to the BLAST score that is generated by sequence comparison of the designated clone with the designated GenBank sequence using the designated BLAST method. This entry is omitted when a clone is included solely due to its association with a particular cluster. If the program used to determine the hit is HMMSW then the score refers to HMMSW score.

P-Value

The entries in the P-Value column refer to the probability that such matches occur by chance.

%Ident

The entries in the “%Ident” column of the table refer to the percentage of identically matched nucleotides (or residues) that exist along the length of that portion of the sequences which is aligned by the BLAST comparison to generate the statistical scores presented. This entry is omitted when a clone is included solely due to its association with a particular cluster.

We claim:

1. A substantially purified nucleic acid molecule that encodes a maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of:

- (a) methionine adenosyltransferase,
- (b) S-adenosyl-methionine decarboxylase,
- (c) aspartate kinase,
- (d) aspartate-semialdehyde dehydrogenase,
- (e) cystathionine gamma-synthase,
- (f) cystathionine beta-lysase, and
- (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase.

2. The substantially purified nucleic acid molecule according to claim 1, wherein said nucleic acid molecule comprises a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

3. A substantially purified maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of

- (a) methionine adenosyltransferase or fragment thereof;
- (b) S-adenosyl-methionine decarboxylase or fragment thereof;
- (c) aspartate kinase or fragment thereof;
- (d) aspartate-semialdehyde dehydrogenase or fragment thereof;
- (e) cystathionine gamma-synthase or fragment thereof;
- (f) cystathionine beta-lysase or fragment thereof; and

(g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof.

4. A substantially purified maize or soybean enzyme or fragment thereof according to claim 3, wherein said maize or soybean enzyme or fragment thereof is encoded by a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

5. A substantially purified antibody or fragment thereof which is capable of specifically binding to a specific maize or soybean enzyme or fragment thereof according to claim 4.

6. A transformed plant having a nucleic acid molecule which comprises:

(A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule;

(B) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of

(a) a nucleic acid sequence which encodes for methionine adenosyltransferase or fragment thereof;

(b) a nucleic acid sequence which encodes for S-adenosyl-methionine decarboxylase or fragment thereof;

(c) a nucleic acid sequence which encodes for aspartate kinase or fragment thereof;

(d) a nucleic acid sequence which encodes for aspartate-semialdehyde dehydrogenase or fragment thereof;

(e) a nucleic acid sequence which encodes for cystathionine gamma-synthase or a fragment thereof;

(f) a nucleic acid sequence which encodes for cystathionine beta-lyase or a fragment thereof;

(g) a nucleic acid sequence which encodes for 5-methyltetrahydropteroyl-triglutamate-homocysteine-S-methyltransferase or a fragment thereof; and

(h) a nucleic acid sequence which is complementary to any of the nucleic acid sequences of (a) through (g); and

(C) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.

7. The transformed plant according to claim 5, wherein said structural gene is complementary to any of the nucleic acid sequences of (a) through (g).

8. A method for determining a level or pattern in a plant cell of an enzyme in a plant metabolic pathway comprising:

(A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence of SEQ ID NO: 1 through SEQ ID NO: 3204 or compliments thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said enzyme;

(B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and

(C) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said enzyme in said plant metabolic pathway.

9. The method of claim 8, wherein said level or pattern is detected by *in situ* hybridization.

10. A method of determining a mutation in a plant whose presence is predictive of a mutation affecting a level or pattern of a protein comprising the steps:

(A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid, said marker nucleic acid selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having a nucleic acid sequence selected from the group of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from said plant, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting said level or pattern of said plant methionine pathway enzyme in said plant;

(B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant; and

(C) detecting the presence of said polymorphism, wherein the detection of said polymorphism is predictive of said mutation.

11. A method of producing a plant containing an overexpressed protein comprising:

(A) transforming said plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein said promoter region is linked to a structural region, wherein said structural region has a nucleic acid sequence selected from group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 wherein said structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of

ABSTRACT

The present invention is in the field of plant biochemistry. More specifically the invention relates to nucleic acid sequences from plant cells, in particular, DNA sequences from maize and soybean plants associated with the methionine pathway. The invention encompasses nucleic acid molecules that encode proteins and fragments of proteins. In addition, the invention also encompasses proteins and fragments of proteins so encoded and antibodies capable of binding these proteins or fragments. The invention also relates to methods of using the nucleic acid molecules, proteins and fragments of proteins and antibodies, for example for genome mapping, gene identification and analysis, plant breeding, preparation of constructs for use in plant gene expression and transgenic plants.

<110> Bledig, Stefan A.
 Byrum, Joseph R.
 Liu, Jingdong
 Hinkle, Gregory J.

<120> Nucleic Acid molecules And Other Molecules Associated With The
 Methionine Synthesis And Degradation Pathways

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<400> 30

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 GTGACAGTGG AGTACCGCAA CGAGGGTGGC GCCATGGTTC CCATCCGTGT GCACACAGTC
 CTCATCTCTA CCCAGCACGA CGAGACAGTC ACCAACGACG AGATTGCTGC TGACCTGAAG
 GAGCACGTCA TCAAGCCAGT CATCCCCGAG CAGTACCTCG ACGAGAAGAC AATCTTCCAC
 CTCAACCCGT CTGGCCGNTT CGTCATCGGC GGACCTNACG GCGACGCCGG CCTACTGGCC
 GNAAGATCAT CATCGACACC TACGGTGGCT GGGGAGCCCA CGGCGGGGGC GCCTTCTTCG
 GCAAGGACCC GACCAANGTG GACCGCACGG GGCCTACGTN CGAGGNAAGC TTGCNA

ctcacagagg ttgcaagaa tggaacctgg cctgggtca ggcccgatgg gaagaccag 60
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 ctcatctcta cccagcacga cgagacagtc accaacgacg agattgctgc tgacctgaag 180
 gagcacgtca tcaagccagt catccccgag cagtacctcg acgagaagac aatcttccac 240
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 gnaagatcat catcgacacc tacgggtggct ggggagccca cggcgggggc gccttcttcg 360
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<210> 31
 <211> 484
 <212> nucleic acid
 <213> Zea mays

<400> 31

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 cgcagcggag cctatgtcgc aaggcaggct gccaaagaca tcgtcgccag cggccttgct 180
 cgccgcgcca tcgtccaggt gtcttacgcc atcggcgtgc ccgagcctct ctccgtgttc 240
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gggccgctac ctnaagacgg cggncctacgg gcacttttgg aaagggacca acctgacttn 420
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 ttaa 484

<210> 32
 <211> 477
 <212> nucleic acid
 <213> Zea mays
 <400> 32

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 ggctcgccc gccgtgcct cgtgcagggtg tcgtacgcca tcggtgtgcc ggagccctgt 180
 ccgtgttcgt cgactcgta cggcaccggca cgatccccga caaggagatc ctcaagtcgt 240
 gaaggagaac ttgcacttca ggcccgggat gatcagcatc aacctcgacc tgaagagggc 300
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 cctgggaagt ggtgaagccc ctcaagtctg acaaggcatc cggcttaang ttggaatnnt 420
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<210> 33
 <211> 488
 <212> nucleic acid
 <213> Zea mays
 <400> 33

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 gtcggtagag cgagaagaag gcaatggcgg ccgagagctt ctttttcacc tcggagtccg 180
 tgaacgaggg gcacccccgac aagctgtgcg accagggtgc ggacgccgtg cttgacgcat 240
 gcctcgcgca ggacccccgac agcaagggtg cctgcgagac ctgcaccaag accaacaatgg 300
 tgatggtggt cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg 360
 acacctgccg cgagatcggg ttacactccg acgacgtggg cctcgacgcc gaccgctgca 420
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cacgaaac

488

<210> 34
<211> 480
<212> nucleic acid
<213> Zea mays

<400> 34

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gaggtcggta gagcgagaag aaggcaatgg cggccgagag cttccttttc acctcggagt 180
ccgtgaacga ggggcacccc gacaagctgt gcgaccaggt gtcggacgcc gtgcttgacg 240
catgcctcgc gcaggacccc gacagcaagg tggcctgcga gacctgcacc aagaccaaca 300
tggtgatggt gttcggcgag atcacgacca aggcgaccgt ggactacgag aagatcgtgc 360
gcgacacctg ccgcgagatc gggttcacct ccgacgacgt gggcctcgac gccgaccgct 420
gcaaggtgct ggtgaacatc gagcagcagt ccccgacat cgcgcanggc gtgcacgggc 480

<210> 35
<211> 453
<212> nucleic acid
<213> Zea mays

<400> 35

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ccgtgaacga ggggcaccca gacaagctgt gcgaccaggt gtcggacgcg gtgctggacg 180
cctgcctggc gcangacccc gacagcaagg tggcctgcna gacctgcacc aagacgaaca 240
tggtgatggt gttcggcgag atcaccacca aggcgagcgt ggactacgag aagatcgtgc 300
gcgacacctg ccgcgagatc gggttcacct tcgacgacgt ggggctcgac nccnaccgct 360
gcaaggtgct ggtgaacatc gagcagcagt ccccgacat cgcgcaaggc gtgcacggca 420
ctttacgaaa ccggcccagag gagatcggcc cnt 453

<210> 36
<211> 505
<212> nucleic acid


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<213>      Zea mays

<400>      36

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gcagcagcgc aagaggttg tagagcgagc gagaagaagg caatggcggc ggagagcttc  180
ctgttcacct cggagtccgt gaacgagggg caccagaca agctgtgcga ccaggtgtcg  240
gacgcggtgc tggacgcctg cctggcgcan gaccccgaca gcaaggtggc ctgcgagacc  300
tgcaccaaga cgaacatggt gatggtgttc ggcgagatca ccaccaaggc gagcgtggac  360
tacgagaaga tcgtgcgcga cacctgccgc gagatcgggt tcacctncca cgacgtgggg  420
ctcgacgccg accgctgcaa ggtgctggtg aacatcgagc ancagtcccc cgacatcgcg  480
cagggcgtgc acgggcactt tacga                                         505

<210>      37
<211>      447
<212>      nucleic acid
<213>      Zea mays

<400>      37

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tggagctggt gaccaggggc acatgttttg gtatgcgact gacgagacct ctgagctgat  120
gccctcagc catgtccttg ccaccaagct tgggtgctcgt ctcacagagg ttcgcaagaa  180
tggaacctgc ccctggctca ggcccgatgg gaagaccag gtgacagtgg agtaccgcaa  240
cgaggggtgg gccatggttc ccatccgtgt gcacacagtc ctcatctcta cccaacacga  300
cgagacangt caccaacgac gaagattgct gctgacctga aaggaacaac gtcatcaaac  360
caagtcatnc ccgaacagta ctttgacga gaagacaatc tttcanctta acccgtctgg  420
nccntttgtc atnngnggac ctnaacg                                         447

<210>      38
<211>      420
<212>      nucleic acid
<213>      Zea mays

<400>      38
  
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<210> 39
 <211> 499
 <212> nucleic acid
 <213> Zea mays
 <400> 39

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 cctcaagatt gtcaaggaga acttcgattt caggcctggc atcatcatca accttgacct 360
 caagaaaggc ggcaacgggc gctacctnaa gacggcggcc tacggcactt tggaaggagc 420
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 agcnggcttt tttttaaaa 499

<210> 40
 <211> 494
 <212> nucleic acid
 <213> Zea mays
 <400> 40

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 gcgagaagaa ggcaatggcg gccgagagct tccttttcac ctcggantcc gtgaacgggg 180

gcaccccgac aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcggcag 240
gaccccgaca gcaangtggc ctgcgagacc tgcaccaaga ccaacatggt gatgggttcg 300
gcgagatcac gaccaaggcg accgtggact acgagaagat cgtgcgcgac acctccgcga 360
gatnnggttc acctccgacg aactgggcct cnaacgccga accgctgcaa ggtctgttga 420
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cgaggagatc ngcg 494

<210> 41
<211> 499
<212> nucleic acid
<213> Zea mays

<400> 41
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gcctnctccg gcaaggaccc caccaagggtg gaccgcagcg gcgcctacgt ggccaggcag 180
gccgccaaga gcatcgtggc cagcggcctc gcccgccgct gcctcgtgca ggtgtcgtac 240
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cccgacaagg agatcctcaa gatcgtgaag gagaacttca acttcaggcc cgggatgatc 360
agcatcaacc tcgacctgaa gaanggcggc aacaggttca tcangaccga cgctacggcc 420
acttcggccg tgacgacgcc gactttacct gggaagtggg gaagcccctt aagttccaca 480
aggattnggt ttaaggttg 499

<210> 42
<211> 325
<212> nucleic acid
<213> Zea mays

<400> 42
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gatcgccgcc gacctcaagg agcacgtcat caagcccgtc atcccggaga ggtacctgga 180
cgagaagacc atcttccacc tcaacccgtc ggggcgcttc gtcacggcg ggcacacgg 240

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cggcgggggc gccttctccg gcaag 325

<210> 43
<211> 319
<212> nucleic acid
<213> Zea mays

<400> 43

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ccgaggtgcg caagaacggc acctgcgcct ggctgaggcc cgacggcaag acccaggtga 180
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acgtcatcaa gcccgtgat 319

<210> 44
<211> 429
<212> nucleic acid
<213> Zea mays

<400> 44

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gagacagtca ccaacgacga gattgctgct gacctgaagg agcacgtcat caagccagtc 180
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nccgcggnc 429

<210> 45
<211> 315
<212> nucleic acid
<213> Zea mays

<400> 45

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cctgcgcctg gctgaggccc gacggcaaga ccaggtgac ggtggagtac gtgaacgagg 180

gcggcgccat ggtgcccgtc cgcgtgcaca ccgtgctcat ctccaccag cacgacgaga 240

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ctgagaagta cctcg 315

<210> 46

<211> 474

<212> nucleic acid

<213> Zea mays

<400> 46

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aaggtggacc gcagcggggc ctacgtcgcg aggcaggctg ccaagagcat cgtcgccgcc 180

ggcctcgccc gccgtgccat cgtccaggtc tctacgcca tcggcgtgcc cgagcccctg 240

tcggtgttcg tggacacgta cggcaccggc gcgatccccg acaaggagat cctgaagatc 300

gtgaaggaga acttcgactt caggcccggc atgatcatca tcaacctcga cctcaagaaa 360

ggcggcaacg ggcgtacct caagacggcg gcctacgggc actttgggag ggacgaaccc 420

gacttcacct gggaagtngt taaaccccc naaggcgga aanccttntt ctgg 474

<210> 47

<211> 410

<212> nucleic acid

<213> Zea mays

<400> 47

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ggcncgcgcc tcaccgaggt tccgcaagac gggnacctgc gcctggntga nggcccgcgc 300
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<210> 48
<211> 297
<212> nucleic acid
<213> Zea mays
<400> 48

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cacgggggacg ccggcctcac ccggcgcgaag atcatcatcg acacctacgg cggctgggga 240
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<210> 49
<211> 438
<212> nucleic acid
<213> Zea mays
<400> 49

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cgatccccga caaggagatc ctgaagatcg tgaaggagaa cttcgacttc angcccggca 300
tgatcatcat caacctngac ctcaagaaag gcggnnacgg nccgtacct taaanaacgg 360
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cccttaaggc ggaaaaaa 438

<210> 50
<211> 316
<212> nucleic acid
<213> Zea mays

<400> 50

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cgacacctgc cgcgagatcg ggttcacctc cgacgacgtg gggctcgcg ccgaccgctg 240
caaggtgctg gtgaacatcg ancagcagtc ccccgacntc ggcgagggcg tgcacgggca 300
nttcacgaag cggccc 316

<210> 51

<211> 339

<212> nucleic acid

<213> Zea mays

<400> 51

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tcttccacct caaccgctct ggccgcttcg tcctcgggcg acctcacggc gacgcggctc 180
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cttctccggc aangacccga ccaaggtgga ccgcagcggg gcctacgtcg cgaggcaggc 300
tgccaagagc atcgtcgccg cggcctcgcc gcngcgctt 339

<210> 52

<211> 300

<212> nucleic acid

<213> Zea mays

<400> 52

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gcgtgcacgg gcacttcacg aagcggcccg aggagatcgg cgcgggacgac cagggccaca 180
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<210> 53

<211> 303
 <212> nucleic acid
 <213> Zea mays

 <400> 53

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 ggagtacgtg aacgagggcg gcgccatggt gcccgccgc gtgcacaccg tgctcatctc 240
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 cat 303

<210> 54
 <211> 477
 <212> nucleic acid
 <213> Zea mays

 <400> 54

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 catcgtgcc gccggcctcg cccgccgcgc cattgtccag gtctcctacg ccacggggtg 180
 cccgagcccc ttctgggtgtt cgtggacacg tacggcaccg gcgcgatccc cgacaagaga 240
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 cctcaagaaa ggoggcaacg ggcgctacct caagacggcg gcctacgggc acttgggagg 360
 gacgaccccg acttcacctg ggaggtggtg aagccctca aggcggaaaa acctttctg 420
 caagaaggcg ccccccggt ttggaanaa gcttttggtc tggctggtc tgtctgg 477

<210> 55
 <211> 487
 <212> nucleic acid
 <213> Zea mays

 <400> 55

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gtcggtagag cgagaagaag gcaatggcgg ccgagagctt ccttttcacc tcggagtccg 180
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tgatgggtgtt cggcgaagat cacgaccaag gcgaccgtgg actacgaaga agatcgtgcg 360
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acttcac 487

<210> 56
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 56
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gctcaggggtg tgcattggcca cttcaccaag cgccccgagg agattggagc tggtgaccag 240
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<210> 57
<211> 315
<212> nucleic acid
<213> Zea mays

<400> 57
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gcgcatgggt tccca 315

<210> 58

<211> 328
 <212> nucleic acid
 <213> Zea mays

 <400> 58

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 tcatcatcga cacctacggt ggctggggag cccacggcgg gggcgccctc tccggcaagg 120
 acccgaccaa ggtggaccgc agcggggcct acgtcgcgag gcaggctgcc aagagcatcg 180
 tcgcccggg cctcgccgc cgtgccatcg tccaggtctc ctacgccatc ggcgtgccc 240
 ancccctgtc ggtgttcgtg gacacgtacg gcaccggcgc gatccccgac aaggagatcc 300
 tgaagatcgt gaaggagaat tcgacttc 328

<210>
 <211>
 <212>
 <213>
 <400>
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 ccgcaaggac
 gtacgtgaac
 ccagcacgac
 caagcccgtc

59
 291
 nucleic acid
 Zea mays

 59

 cgagctgatg ccgctgagcc acgtgctggc caccaagctc ggcgcgcgnc tgacggaggt 60
 ccgcaaggac ggcacctgcn cctggctcag gcccgacggc aagaccagagg tgacggtgga 120
 gtacgtgaac ganggcggcg ccatggtgcc cgtccgctg cacaccgtgc tcatctccac 180
 ccagcacgac gagaccgtca ccaacgacga gatcgccgcc gacctcaagg agcacgtcat 240
 caagcccgtc atcccggaga ggtacctgga cgagaagacc atcttccacc t 291

<210> 60
 <211> 305
 <212> nucleic acid
 <213> Zea mays

 <400> 60

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 ctaggtgctc gtctcaccga ggtccgcaag aacggaacct gccctggct caggcctgat 120
 gggaagaccc aggtgacagt cgagtaccgc aatgaggggtg gtgccatggt ccccatccgt 180
 gtccacaccg tctcatctc caccagcac gacgagacag tgaccaatga tgagatcgct 240
 gctgacctga aggagcatgt catcaagcct gtcacccctg agcagtacct tgacgagaag 300

accat

305

<210> 61
<211> 300
<212> nucleic acid
<213> Zea mays

<400> 61

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ggcgaccgtg gactacgaga agatcgtgcg cgacacctgc cgcgagatcg ggttcacctc 120
cgncgacgtg ggctcgacg ccgncgcgtg caaggtgctg gtgaacatcg agcagcagtc 180
ccccgacatc gcgcagggcg tgcacgggca cttcacgnag cggnccgagg agatnggngc 240
gggcgaccag ggncacatgt tcgggtacgn caccgacgag acccccgagc tgatgccgct 300

gagcagcagc gcaagaggtc ggtagagcga gaagaaggca atggcggccg agagcttcct

<210> 62
<211> 558
<212> nucleic acid
<213> Zea mays

<400> 62

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ggaattcccg ggtcgaccca cgcgtccgcc cagcgtccg cccacgcgtc cgcccacgog 120
tccgcccacg cgtccgccc cgcgtccggc ctcttctccc tcttgccggt cccgaataaa 180
gagcagcagc gcaagaggtc ggtagagcga gaagaaggca atggcggccg agagcttcct 240
tttcacctcg gagtccgtga acgaggggca ccccgacaag ctgtgcgacc aggtgtcgga 300
cgccgtgctt gacgcatgcc tcgcgagga ccccgacagc aaggtggcct gcgagacctg 360
caccaagacc aacatggtga tgggtgttcg cgagatcacg accaaggcga ccgtggacta 420
cgagaagatc gtgcgagaca cctgccgca gatcgggttc acctncgacg acgtgggcct 480
tnacgcenac cnntgcaagg tncgtgtgaa cattgagcaa naattcccng gactttnnge 540
anggcgttca cnggcant 558

<210> 63
<211> 332
<212> nucleic acid
<213> Zea mays

<400> 63

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ggtgccatgg tccccatccg tgtccacacc gtctcatctt ccaccagca cgacgagaca 120

gtgaccaatg atgagatcgc tgctgacctg aaggagcatg tcatcaagcc tgtcatccct 180

gagcagtacc ttgacgagaa gaccatcttc caccttaacc catctggccg ctttgtcatg 240

tggacctcac ggcgatgctg gcctcactgg ccgcaagatc atcatgacac ctacgggtggc 300

tggggagccc atggtggtgg cgctttctcc gg 332

<210> 64

<211> 314

<212> nucleic acid

<213> Zea mays

<400> 64

ctcggagtct gtgaacgagg gacaccctga caagctctgt gaccaggtct cagatgccgt 60

tcttgacgct tgccttgctg aggaccctga cagcaagggt gcttgtgaga cctgcaccaa 120

gaccaacatg gtcatggtct ttggtgagat caccaccaag gccaatgttg actacgagaa 180

gattgtcagg gagacctgcc gcaacattgg ttttgtgtca aacgatgttg ggcttgacgc 240

cgaccactgc aagtgtctgt gaacattnag cagcagtcct ctgatattgc tcanggtgtg 300

catggccact tcac 314

<210> 65

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 65

ggacgagaag accatcttcc acctcaaccc gtcggggcgc ttctcatcg gcgggcccc 60

cggggacgcc ggctcaccg gccgcaagat catcatcgac acctacggcg gctggggagc 120

ccacggcggg ggcgcttct ccggcaagga cccaccaag gtggaccgca gcggggccta 180

cgtcgccagg caggccgcca agagcatcgt ggccagcggc ttgcccgcg gctgcctcgt 240

gcaggtgtcc tacgccatcg ggtgccggag cccctgtccg tgttcgtcga ctc 293

<210> 66

<211> 289
 <212> nucleic acid
 <213> Zea mays

 <400> 66

 gcaaggtggc ctgcgagacc tgcaccaaga cgaacatggt gatggtgttc ggcgagatca 60
 ccaccaaggc gagcgtggan tacgagaagn tcgtgcgcga cacctgccgc gagatcgggt 120
 tcacctccga cgacgtgggg ctgcacgccg accgctgcaa ggtgctggtg aacatcgagc 180
 agcagtcccc cgacatcgcg cagggcgtgc acgggcactt cacgaagcgg cccgaggaga 240
 tcggcgccgg cgaccagggc cacatgttcg ggtacgccac cgacgagac 289

<210> 67
 <211> 306
 <212> nucleic acid
 <213> Zea mays

 <400> 67

 gttgactacg agaagattgt caggagagacc tgccgcaaca ttggttttgt gtcaaacgat 60
 gttgggcttg acgccgacca ctgcaagggtg ctctgaaca ttgagcagca gtccccgat 120
 attgctcagg gtgtgcatgg ccacttcacc aagcgccccg aggagattgg agctggtgac 180
 caggacaca tgttcgggta tgcgaccgat gagaccctg agttgatgcc cctcagccat 240
 gtcttgcca ccaagctagg tgctcgtctc accgaggtcc gcaagaacgg aacctgcccc 300
 tggctc 306

<210> 68
 <211> 303
 <212> nucleic acid
 <213> Zea mays

 <400> 68

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 tgacagcaag gttgcttggtg agacctgcac caagaccaac atggtcatgg tctttggtga 120
 gatcaccacc aaggccaatg ttgactacga gaagattgtc agggagacct gccgcaacat 180
 tggttttgtg tcaaacgatg ttgggcttga cgccgaccac tgcaagggtc tcgtgaacat 240
 tgagcagcag tccccgata ttgctcaggg tngcatggc cacttcacca agcgccccga 300

gga

303

<210> 69
<211> 300
<212> nucleic acid
<213> Zea mays

<400> 69

caaagaccaa catggtcatg gtcttttggtg agatcaccac caaggccaat gtcgactacg 60
agaagattgt cagggagaca tgccgcaaca ttggtttcgt ntcgaacgat gtcgggcttg 120
acgtgacca ctgcaagggtg cttgtgaaca ttgagcagca gtcccctgat attgctcagg 180
gtgtncacgg ccacttcacc aagcgccccg aggagattgg agctggtgac caggggcaca 240
tgtttgggta tgcgactgac gagaccctg agctgatgcc cctcagccat gtccttgcca 300

<210> 70
<211> 329
<212> nucleic acid
<213> Zea mays

<400> 70

gatcaaagaa gatggcagct gtcgacacat tcctcttcac ctcgaggtct gtgaacgagg 60
gacaccctga caagctctgt gaccaggtct cagatgccgt tcttgacgct tgccttgctg 120
aggaccctga cagcaagggtt gcttgtgaga cctgcaccaa gaccaacatg gtcattggtct 180
ttggtgagat caccaccaag gccaatgttg actacgagaa gattgtcagg gagacctgcc 240
gcaacattgg ttttgtgtca aacgatgttg ggcttgacgc cgaccattgc aagggtgcncg 300
tgaanatnng cancagtcct ctgatattg 329

<210> 71
<211> 304
<212> nucleic acid
<213> Zea mays

<400> 71

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gtcttttgng agatcaccac caaggccaat gtcgactacg agaagattgt cagggagaca 120
tgccgcaaca ttggtttcgt gtcgaacgan gtcgggcntg angctgacca ctgcaaggng 180

cttgtgaaca ttgagcagca gtccccctgat attgctcagg gtgtgcacgg ccacttcacc 240
aagcgccccg aggagattgg agctggtgac caggggcaca tgtttgggta tgcgactnac 300
gaga 304

<210> 72
<211> 307
<212> nucleic acid
<213> Zea mays

<400> 72

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catggtcang gnccttttggg gagatcacca ccaaggccaa tgttgactac gagaagattg 120
tcagggagac ctgccgcaac attggttttt tgtcaaacga tgttgggctt gacgccgacc 180
actgcaaggt gctcgtgaac attgagcagc agtccccctga tattgctcag ggtgtgcatg 240
gccacttcac caagcgcccc gaggagattg gagctggtga ccagggacac atgttcgggt 300
atgcgac 307

<210> 73
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 73

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gcgcgcctga cggaggtccg caaggacggc acctgcgcct ggctcaggcc cgacggcaag 120
accaggtga cgggtggagta cgtgaacgag ggcgggcgcca tgggtgcccgt ccgcgtgcac 180
accgtgctca tctccaccca gcacgacgag accgtcacca acgacgagat cgccgccgat 240
ctcaaggagc acgtcatcaa gcccgtcac ccggagaggt ac 282

<210> 74
<211> 320
<212> nucleic acid
<213> Zea mays

<400> 74

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cgatgggaag acccaggtga cagtggagta ccgcaacgag ggtggcgcca tggttcccat 120
ccgtgtgcac acagtcctca tctctacca gcacgacgag acagtcacca acgacgagat 180
tgctgctgac ctgaaggagc acgtcatcaa gccagtcac cccgagcagt acctcgacga 240
gaagacaatc ttccacctca acccgtctgg ccgcttcgtc atcggcggac tcacggcgac 300
ctggcctcac tggccggaag 320

<210> 75
<211> 370
<212> nucleic acid
<213> Zea mays
<400> 75

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aggagatcct caagatcgtc aaggagaact tcgacttcag gcccgggatg atcaccatca 180
acctcgacct caagaagggc ggcaacaggt tcatcaagac cgccgcatac ggccactttg 240
gccgtgacga cgccgaactc acctgggagg tggtaagcc cctaaagaag gcatccgctt 300
aagaatgtat tgggaagttc actggacatg aggttcatct tcgtctggct ctgctgatac 360
ctgcaaggat 370

<210> 76
<211> 300
<212> nucleic acid
<213> Zea mays
<400> 76

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acgaggggtg cgccttggtt cccatccgtg tgcacacagt cctcatctct acccagcacg 120
acgagacagt caccaacgac gagattgctg ctgacctgaa ggagcacgtc atcaagccag 180
tcatccccga gcagtacctc gacgagaaga caatcttcca cctcaacccg tctggccgct 240
tcgtcatcgg cggacctcac ggcgacgtg gcctcactgg ccggaagatc atcatcgaca 300

<210> 77
<211> 315
<212> nucleic acid

<213> Zea mays

<400> 77

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gacctgcacc aagaccaaca tggatcatggt ctttggtgag atcaccacca aggccaatgt 120
tgactacgag aagattgtca gggagacctg ccgcaacatt ggttttgtgt caaacgatgt 180
tgggcttgac gccgaccact gcaaggtgct cgtgaacatt gagcagcagt cccctgatat 240
tgctcagggt gtgcatggcc attcaccaag cgccccgang agattggagc tggtgaccag 300
gacacatggt cgggg 315

<210> 78

<211> 297

<212> nucleic acid

<213> Zea mays

<400> 78

ctcttcacct cggagtctgt gaacgagggg caccctgaca agctctgtga ccaggctctca 60
gatgccgttc ttgacgcttg ccttgctgag gaccctgaca gcaaggttgc ttgtgagacc 120
tgcaccaaga ccaacatggt catggtcttt ggtgagatca ccaccaaggc caatgttgac 180
tacgagaaga ttgtcagggg gacctgccgc aacattgggt ttgtgtcaaa cgatgttggg 240
cttgacgcgc accactgcaa gtgctcgtga acattgagca gcagtcccct gatattg 297

<210> 79

<211> 448

<212> nucleic acid

<213> Zea mays

<400> 79

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gnacattgg ttttgtgtca aacgatgttg ggcttgacgc cgaccactgc aagggtgctcg 120
tgaacattna gcagnagtnc cctgatattg ctanggtgt gcatggccac ttnaccaanc 180
gccccganga gattgganct ggtgaccagg gacacatggt cgggtatgcy accgatgaga 240
cccctnagtt gatgccctc agccatgtcc ttgccaccaa gctaggtgct cgtctnaccg 300
aggtncncaa gaaccggaac ctgccnctgg ctangcctg atgngaagac cnatgtgaca 360

gtcnantnnc gnaatgaagg gtggtgccat tgncccac ctngtcaaca ccgttcttat 420
 ttcaaccaag tnngacgagg acaatgac 448

<210> 80
 <211> 287
 <212> nucleic acid
 <213> Zea mays
 <400> 80

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 cctgaaggag catgtcatca agcctgtcat ccctgagcag taccttgacg agaagaccat 120
 cttccacctt aacccatctg gccgctttgt cattgggtgga cctcacggcg atgctggcct 180
 cactggccgc aagatcatca ttgacaccta cgggtggctgg ggagcccatg gtggtggcgc 240
 tttctccggc aaggacccaa ccaaggttga ccgcagcgga gctatgt 287

<210> 81
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 81

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 acgacgagac agtgaccaat gatgagatcg ctgctgacct gaaggagcat gtcacaaagc 120
 ctgtcatccc tgagcagtac cttgacgaga agaccatctt ccaccttaac ccatctggcc 180
 gctttgtcat tgggtggacct cacggcgatg ctggcctcac tggccgcaag atcatcattg 240
 acacctacgg tggctgggga gcccatggtg gtggcgcttt ctccggcaag 290

<210> 82
 <211> 287
 <212> nucleic acid
 <213> Zea mays
 <400> 82

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 cgaacatggt gatggtgttc ggcgagatca ccaccaaggc gagcgtggac tacgagaaga 120
 tcgtgcgcga cacctgccgc gagatcgggn nnacctccga cgacgtgggg ctgcagcccg 180

accgctgcaa ggtgctggtg aacatcgagc agcagtcgcc cgacatcgcg cagggcgtgc 240
acgggcactt cacgaagcgg cccgaggaga tcggcgccgg cgaccag 287

<210> 83
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 83

atgcccctca gccatgtcct tgccaccaag ctaggtgctc gtctcaccga ggtccgcaag 60
aacggaacct gccctggct caggcctgat gggaagacct aggtgacagt cgagtaccgc 120
aatgaggggtg gtgccatggt ccccatccgt gtccacaccg tcctcatctc caccagcac 180
gacgagacag tgaccaatga tgagatcgct gctgacctga aggagcatgt catcaagcct 240
gtcatccctg agcagtacct tgacgagaag accatcttcc accttaacct a 291

<210> 84
<211> 283
<212> nucleic acid
<213> Zea mays

<400> 84

tgccgctgag ccacgtgctg gccaccaagc tgggcgcgcg cctcaccgag gtgcgcaaga 60
acggcacctg cgcctggctg aggcccgatc ggcaagacct aggtgacggt ggagtacgtg 120
aacgagggcg gcgccatggt gcccgctccg gtgcacaccg tgctcatctc caccagcac 180
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gtgatccctg agaagtacct cgacgagaag accatcttcc acc 283

<210> 85
<211> 274
<212> nucleic acid
<213> Zea mays

<400> 85

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atgcctcgcg caggaccccg acagcaaggt ggctgcgag acctgcacca agaccaacat 120
ggtgatgggtg ttcggcgaga tcacgaccaa ggcgaccgtg gactacgaga agatcgctgcg 180

cgacacctgc cgcgagatcg ggttcacctc cgacgacgtg ggctcgcaca ccgaccgctg 240
 caaggtgctg gtgaacatcg agcagcagtc cccc 274

<210> 86
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 86

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 ccggaagatc atcatcgaca cctacgggtgg ctgggggagcc cacggcgggg ggcgccttctc 180
 cggcaaggac ccgaccaagg tggaccgcag cgggggctac gtcgcgaggc aggttgccaa 240
 gagcatcgtc gccgcgggcc tcgccgcng tgccatcgtc caggtctcct 290

<210> 87
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 87

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 gaccaggctc cagatgccgt tcttgacgct tgccttgctg aggaccctga cagcaagggtt 120
 gcttgtgaga cctgcaccaa gaccaacatg gtcattgtct ttggtgagat caccaccaag 180
 gccaatgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240
 aacgatgttg ggcttgacgc cgaccactgc aaggtgctcg tgaacattga 290

<210> 88
 <211> 288
 <212> nucleic acid
 <213> Zea mays
 <400> 88

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 tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc cgacagcaag 120
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acaccgtcct catctccacc cagcacgacg agacagtgac caatgatgag atcgctgctg 180
acctgaagga gcatgtcatc aagcctgtca tccctgagca gtaccttgac gagaagacca 240
tcttccacct taacccatct ggccgctttg tcattggtgg acctcacggc g 291

<210> 92
<211> 285
<212> nucleic acid
<213> Zea mays

<400> 92

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cccctgtccg tgttcgtcga ctctacggc accgggacca tccccgacaa ggagatccta 120
aagatcgtca aggagaactt cgacttcagg ccagggatga tcaccatcaa cctcgacctc 180
aagaagggcg gcaacagggt catcaagacc gccgcatacg gccactttgg ccgtgacgac 240
gccgacttca cctgggagggt ggtcaagccc ctaaagaagg catcc 285

<210> 93
<211> 283
<212> nucleic acid
<213> Zea mays

<400> 93

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tatgcgaccg atgagacccc tgagttgatg ccctcagcc atgtccttgc caccaagcta 120
ggtgctcgtc tcaccgaggt ccgcaagaac ggaacctgcc cctggctcag gcctgatggg 180
aagaccaggg tgacagtcga gtaccgcaat gaggggtggg ccatggtccc catccgtgtc 240
cacaccgtcc tcctctccac ccagcacgac gagacagtga cca 283

<210> 94
<211> 298
<212> nucleic acid
<213> Zea mays

<400> 94

actacgagaa gattgtcagg gagacatgcc gcaacattgg ttctgtgtcg aacgatgtcg 60
ggcttgacgc tgaccactgc aaggtgcttg tgaacattga gcagcagtc cctgatattg 120

ctcaggggtgt gcacggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 180
 ggcacatggt tgggtatgcn actgacgaga cccttgagct gatgcccctc agccatgtcc 240
 ttgccaccaa gcttgggtgtc gtctcacnga aggttcgcaa gaatggaacc tgcccctt 298

<210> 95
 <211> 469
 <212> nucleic acid
 <213> Zea mays

<400> 95

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 ccgacaagct gtgcnaccag gtgtcggacg ccgtgcttga cncatgcctc gcgcaggacc 180
 ccnacagcaa ggtggcctgc nagacctgca ccaanaccaa catggtgatg gtgttcggcg 240
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 atcgggttca ccttcgncga cgtgngccct tgactccnnc ccggtgcaag gtgctggtga 360
 acattnatca tcaatncccc gacattnttc aaggcnttca cggcacttta cgaaacggcc 420
 cnangagatc ggccggggcca acagngccac atnttcgggt ccccccca 469

<210> 96
 <211> 293
 <212> nucleic acid
 <213> Zea mays

<400> 96

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 cctgattgct caggggtgtgc atggccactt caccaagcgc cccgaggaga ttggagctgg 120
 tgaccaggga cacatgttcg ggtatgcnac cgatgagacc cctgagttga tgcccctcag 180
 ccatgtcctt gccaccaagc taggtgctcg tctcaccgag gtccgcaaga acggaacctg 240
 cccctggctc aggctgatg ggaagacca ggtgacagtc gagtaccgca aaa 293

<210> 97
 <211> 280
 <212> nucleic acid
 <213> Zea mays

<400> 97

cggnacgntg gcgtgaacga ggggcaccca gacaagctgt gcgaccaggt gtcggacgcg 60

gtgctggacg cctgcctggc gcagganccc gacagcaagg tggcctgcga gacctgcacc 120

aagacgaaca tggatgatggg gttcggcgag atcaccacca aggcgagcgt ggactacgag 180

aagatcgtgc gcgacacctg ccgcgagatc gggttcacct ccgacgacgt ggggctcgac 240

gccgaccgct gcaagggtgct ggtgaacatc gagcagcagt 280

<210> 98

<211> 285

<212> nucleic acid

<213> Zea mays

<400> 98

catggtggtg gcgctttctc cggcaaggac ccaaccaagg ttgaccgcag cggagcctat 60

gtcgcgaggc aggctgccaa gagcatcgtc gccagcggcc ttgctcgccg cgccatcgtc 120

cagggtgtctt acgccatcgg cgtgcccagag cctctctccg tgttcgtcga cacgtacggc 180

accggcgcga tccccgacaa ggagatcctc aagattgtca aggagaactt cgatttcagg 240

cctggcatga tcatcatcaa ccttgacctc aagaaaggcg gcaag 285

<210> 99

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 99

aggtgacagt cgagtaccgc aatgaggggtg gtgccatggt ccccatncgt gtccacaccg 60

tcctcatctc caccagcac gacgagacag tgaccaatga tgagatcgct gctgacctga 120

aggagcatgt catcaagcct gtcatncctg agcagtacct tgacgagaag accatcttcc 180

accttaacct atctggccgc tttgtcattg gtggacctca cggcgatgct ggcctcactg 240

gccgcaagat catcattgac acctacggtg gctgggga 278

<210> 100

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 100

gtgaccaatg atgagatcgc tgctgacctg aaggagcatg tcatcaagcc tgtcatccct 60

gagcagtacc ttgacgagaa gaccatcttc caccttaacc catctggccg ctttgcatt 120

ggtggacctc acggcgatgc tggcctcact ggccgcaaga tcatcattga cacctacggt 180

ggctggggag cccatggtgg tggcgctttc tccggcaagg acccaaccaa ggttgaccgc 240

agcggancct atgtcgcaag gcaggctgcc aagag 275

<210> 101

<211> 291

<212> nucleic acid

<213> Zea mays

<400> 101

gatcgctgct gacctgaagg agcatgtcat caagcctgtc atccctgagc agtacottga 60

cgagaagacc atcttccacc ttaaccctac tggccgcttt gtcattggtg gacctcacgg 120

cgatgctggc ctactggcc gcaagatcat cattgacacc tacggtggct ggggagccca 180

tggtggtggc gctttctccg gcaaggaccc aaccaagggt gaccgcagcg gaggcctatgt 240

cgcaaggcag gctgccaaga gcatcgtcgc cagcggcttg ctgcgcgcgc c 291

<210> 102

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 102

agaagatggc cggactcgac accttccctc tcacctcgga gtccgtgaac gagggacacc 60

ctgacaagct ctgcgaccag gtctcagatg ctgttctgga cgcttgccctg ctgaggaccc 120

tgacagcaag gttgcttgcg agacctgcac caagaccaac atgggtcatgg tctttggtga 180

gatcaccacc aaggccaatg tcgactacga gaagattgtc agggagacat gccgcaacat 240

tggtttctgt tcgaacgatg tcgggcttga cgctgaccac tgcaagtgt tgtgaacatt 300

g 301

<210> 103

<211> 336

<212> nucleic acid

<213> Zea mays

<400> 103

ctccctcttg ccggtcccga ataaagagca gcagcgcaag aggtcggtag agcgagaaga 60
aggcaatggc ggccgagagc ttctttttna cctcggagtc cgtgaaacga ggggcncccc 120
gacaagctgt gcgaccaggt gtcggacgcc gtgcttgacg catgcctcgc gcaggacccc 180
gacagcaagg tggcctgcga gacctgcacc aagaccaaca tggatgatggt gttcggcgag 240
atcacgacca aggcgaccgt ggactacgag aagatcgtgc gcgacacctg ccgcgagatc 300
gggttcacct ccgacgacgt gggcctcgac gccgac 336

<210> 104

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 104

tgagaagtac ctgcagcaga agaccatctt ccacctcaac ccgtccgggc gcttcgtcat 60
cggcggggccc cagcgtgacg ccggcctcac cggccgcaag atcatcatcg acacgtacgg 120
cggctgggga gcccacggcg gtggcgccctt ctccggcaag gacccacca aggtggaccg 180
cagcggcgcc tacgtggcca ggcaggccgc caagagcatc gtggccagcg gctcgcccgc 240
cgctgcctcg tgcagtgtcg tacgccatcg ctgccg 276

<210> 105

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 105

gagattggag ctggtgacca gggacacatg ttcgggtatg cgaccgatga gacccttgag 60
ttgatgcccc tcagccatgt ccttgccacc aagctaggtg ctggtctcac cgaggctcgc 120
aagaacggaa cctgcccctg gctcaggcct gatgggaaga ccaggtgac agtcgagtac 180
cgcaatgagg gtggtgccat ggtccccatc cgtgtccaca ccgtcctcat ctccaccag 240
cacgacgaga cagtgaccaa tgatgagatc gtgctgacct gaaggag 287

<210> 106

<211> 303
 <212> nucleic acid
 <213> Zea mays

 <400> 106

 accgtcctca tctccaccca gcacgacgag acagtgacca atgatgagat cgctgctgac 60
 ctgaaggagc atgtcatcaa gcctgtcatc cctgagcagt accttgacga gaagaccatc 120
 ttccacctta acccatctgg ccgctttgtc attggtggac ctcacggcga tgctggcctc 180
 actggccgca agatcatcat tgacacctac ggtggctggg gagcccatgg tgggtggcgt 240
 ttctccggca aggacccaac caagttgacc gcagcgganc tatgtcgcaa ggcagctgcc 300
 aag 303

<210> 107
 <211> 279
 <212> nucleic acid
 <213> Zea mays

 <400> 107

 gtgaccaggt ctcagatgcc gttcttgacg cttgccttgc tgaggaccct gacagcaagg 60
 ttgcttgtga gacctgcacc aagaccaaca tggtcatggt ctttggtgag atcaccacca 120
 aggccaatgt tgactacgag aagattgtna gggagacctg ccgcaacatt ggttttgtgt 180
 caaacgatgt tgggcttgac gccgaccact gcaagggtgt cgtgaacatt gagcagcagt 240
 cccctgatat tgctcagggg gtgcatggcc acttcacca 279

<210> 108
 <211> 330
 <212> nucleic acid
 <213> Zea mays

 <400> 108

 ctotttctccc tcttgccggt cccgaataaa gagcagcagc gcaagaggtc ggtagagcga 60
 gaagaaggca atggcggccg agagcttcct ttccacctcg gagtccgtga acgaggggca 120
 ccccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc tcgcgcagga 180
 ccccgacagc aagggtggcct gcgagacctg caccaagacc aacatgggtga tgggtgttcgg 240
 cgagatcacg accaaggcga ccgtggacta cgagaagatc gtgcgcgaca cctgccgcga 300

gatcgggttc acctccgacg acgtgggcct

330

<210> 109
<211> 298
<212> nucleic acid
<213> Zea mays

<400> 109

ccgacggcaa gacccaggtg acggtggagt acgtgaacna nggcggcgcc atggtgcccg 60
tccgntgca caccgtgctc atctccaccc agcacgacga gaccgtcacc aangacgagn 120
tcgcccgcgn cctcanggag cacgtcntna agcccgtcat cccgganagg tacctggacg 180
anaagacnt cttncacctc aaccgctcgg gggegnntcg tentcggcgg gccccacggg 240
gacnccggcc tnaccggccg caagntgntc ntngncacct acngcggntg gggagccc 298

<210> 110
<211> 498
<212> nucleic acid
<213> Zea mays

<400> 110

ccctcttttn gcctatccgg gccgaccac gcgncacgcg gnggctcngn gcgtatcgag 60
cccacggatt ttggntctn ctccggcaag gacccaccn nggtgggggn gnattgggnc 120
ctaccgtcgc caggcangcc gacaagagca tngnggccag cggcctcgn cgcgntgcc 180
tcngncaggt gtctacgcc atcggcgtgc cggagcccct gtccgtgttc gtngactcct 240
acggcaccgg gaccatcccc gacaaggaga tcctaaagat cgtnaaggag aacttcgact 300
tcaggccagg gatggtcacc atcaacctcg acctcaagaa gggcggcaac aggttcatca 360
agaccgcgn atacggccac tttggcccgt gacgacgccg acttcacctg ggaggtggtc 420
aagcccctaa agaaggcatc cgcttaagaa tgtattnga aagttcactg gacatgaagg 480
atcatcttcc tctnggct 498

<210> 111
<211> 284
<212> nucleic acid
<213> Zea mays

<400> 111

gccgcagat caaagaagat ggcagctgtc gacacattcc tttcacctc ggagtctgtg 60
 aacgagggac accctgacaa gctctgtgac caggtctcag atgccgttct tgacgcttgc 120
 cttgctgagg accctgacag caaggttgct tgtgagacct gcaccaagac caacatggtc 180
 atggtctttg gtgagatcac caccaaggcc aatgttgact acgagaagat tgtcaggagg 240
 acctgccgca acattggttt tgtgtcaaac gatgttgggc ttga 284

<210> 112
 <211> 328
 <212> nucleic acid
 <213> Zea mays
 <400> 112

ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga cacctgccgc 60
 gagatcgggt tcacctccga cgacgtgggc ctgcacgccg accgctgcaa ggtgctggtg 120
 aacatcgagc agcagtcctc cgacatcgcg cagggcgctgc acgggcactt cacgaagcgg 180
 cccgaggaga tcggcgcggg nnaccagggc cacatgttcg ggtacgccac cgacgagacc 240
 cccgagctga tgccgctgag ccaacgtgct ggccaacaag ctgggcgcgcg ggctcaccga 300
 ngtgcgaaaa acggcaactg cgctggct 328

<210> 113
 <211> 287
 <212> nucleic acid
 <213> Zea mays
 <400> 113

gggggcgctt tctccggcaa ggacccgacc aaggtggacc gcagcggggc ctacgtcgcg 60
 aggcaggctg ccaagagcat cgtcgccgcc ggctcgcgcc gccgcgccat tgtccaggtc 120
 tctacgcca tcggcgctgc cgagccctt tcggtgttcg tggacacgta cggcaccggc 180
 gccatccccg acaaggagat cctgaagatc gtgaaggaga acttcgactt caggccccgc 240
 atgatcatca tcaacctcga cctcaagaaa ggccgcaacg ggcgcta 287

<210> 114
 <211> 261
 <212> nucleic acid
 <213> Zea mays

<400> 114

cgacgccgaa cgcgtgcaag gtgctggtga acatcgagca gcagtcccc gacatcgcg 60
agggcgtgca cgggcacttc acgaagcggc ccgaggagat cggcgcgggc gaccagggcc 120
acatgttcgg gtacgccacc gacgagaccc ccgagctgat gccgtgagc cacgtgctgg 180
ccaccaagct gggcgcgcg ctcaccgagg tgcgcaagaa cggcacctgc gcctggctga 240
ggcccgcagg caagaccag g 261

<210> 115

<211> 294

<212> nucleic acid

<213> Zea mays

<400> 115

gggccacttc accaagcgcc ccgaggagat tggagctggt gaccagggac acatgttcgg 60
gtatgcgacc gatgagaccc ctgagttgat gccctcagc catgtccttg ccaccaagct 120
aggtgctcgt ctcaccgagg tccgcaagaa cggaanctgc ccctggctca ggctgatgg 180
gaagaccag gtgacagtcg agtaccgcaa tgagggtggt gccatgggcc ccatccgtgt 240
ccacaccgtc ctcatctcca cccagcacga cgagacatga ccaatgatga gata 294

<210> 116

<211> 318

<212> nucleic acid

<213> Zea mays

<400> 116

ctgaaggagc acgtcatcaa gccagtcac cccgagcagt acctcgacga gaagacaatc 60
ttccacctca acccgtctgg ncgcttcgtc atcggcgagc ctcacggcga cgccggcctc 120
actggccgga agatcatcat cgacacctac ggtggctggg gagccacgg cgggggccc 180
ttctccggca aggacccgac caangtgagc cgcagcgggg cctacgtgc gangcaggct 240
gccaagagca tcgtcgccgc cggcctcgcc gcngcgccat cgtccaggtc tctagcatgg 300
gtgccgancc tatcgtgt 318

<210> 117

<211> 256

<212> nucleic acid

<213> Zea mays

<400> 117

gagaagtacc tcgacgagaa gaccatcttc cacctcaacc cgtccgggcg cttcgtcatc 60
 ggcgggcccc acggtgacgc cggcctcacc ggccgcaaga tcatcatcga cacgtacggc 120
 ggctggggag cccacggcgg tggcgcttc tccggcaagg accccaccaa ggtggaccgc 180
 agcggcgctt acgtggccag gcaggccgcc aagagcatcg tggccagcgg cttcgcccg 240
 cgctgccttc tgcaag 256

<210> 118

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 118

gtcgacacat tcctcttcac ctgggagtct gtgaacgagg gacaccctga caagctctgt 60
 gaccaggtct cagatgccgt tcttgacgct tgccttgctg aggaccctga cagcaagggt 120
 gcttgtgaga cctgcaccaa gaccaacatg gtcattgtct ttggtgagat caccaccaag 180
 gncnatgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240
 aacgatgttg ggcttgacgc cgaccactgc aaggt 275

<210> 119

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 119

gtcgacacat tcctcttcac ctgggagtct gtgaacgagg gacaccctga caagctctgt 60
 gaccaggtct cagatgccgt tcttgacgct tgccttgctg aggaccctga cagcaagggt 120
 gcttgtgaga cctgcaccaa gaccaacatg gtcattgtct ttggtgagat caccaccaag 180
 gccattgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240
 aacgatgttg ggcntgacgc cgaccactgc aaggtg 276

<210> 120

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 120

cggcacggtg gtcgcagcat cgctgctgac ctgaaggagc atgtcatcaa gcctgtnatc 60
cctgagcagt accttgacga gaagaccatn ttccacctta acccatctgg ccgctttgtc 120
attggtggac ctcacggcga tgctggcctc actggccgca agatcatcat tgacacctac 180
ggtaggtggg gagcccatgg tggtaggcgt ttctccggca aggacccaac caaggttgac 240
cgcagcggag cctatgtcgc aangcangct gccaaagaca tcgtcgccaa cggcttgctc 300
gccgcgcca 309

<210> 121

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 121

ctcagggtgt gcatggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 60
gacacatgtt cgggtatgcg accgatgaga ccctgagtt gatgcccctc agccatgtcc 120
ttgccaccaa gctaggtgct cgtctcaccg aggtccgcaa gaacggaacc tgcccctggc 180
tcaggcctga tgggaagacc caggtgacag tcgagtaccg caatgagggt ggtgccatgg 240
tccccatccg tgtccacacc gtctctca 267

<210> 122

<211> 277

<212> nucleic acid

<213> Zea mays

<400> 122

gcaaggtgct tgtgaacatt gagcagcagt ccctgatat tgctcagggt gtgcacggcc 60
acttcaccaa gcgccccgag gagattggag ctggtgacca ggggcacatg tttgggtatg 120
cgactgacga gaccctgag ctgatgcccc tcagccatgt ccttgccacc aagcttggtg 180
ctcgtctcac ggaggttcgc aagaatgaa cctgcccctg gtcaggccc gatgggaaga 240
cccagtgaca attggantac cgcaacgagg gtggccc 277

<210> 123

<211> 264
 <212> nucleic acid
 <213> Zea mays

 <400> 123

 gccacatggt cgggtacgcc accgacgaga cccccgagct gatgccgctg agccacgtgc 60
 tggccacca gctggggcgcg cgcctcaccg aggtgcgcaa gaacggcacc tgcgcctggc 120
 tgaggccccga cggcaagacc caggtgacgg tggagtacgt gaacgagggc ggcgccatgg 180
 tgcccgtccg cgtgcacacc gtgctcatct ccaccagca cgacgagacc gtcaccaacg 240
 acgagatcgc ccgccgacct caag 264

<210> 124
 <211> 269
 <212> nucleic acid
 <213> Zea mays

 <400> 124

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 cgccgaccac tgcaaggtgc tcgtgaacat tgagcagcag tcccctgata ttgctcaggg 120
 tgtgcatggc cacttcacca agcgccccga ggagattgga gctggtgacc agggacacat 180
 gttcgggtat gcgaccgatg agaccctga gttgatgccc ctcagccatg tccttgccac 240
 caagctaggt gctcgtctca ccgaggtcc 269

<210> 125
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 125

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 aacggcacct gcgcctggct gaggcccgac ggcaagacct aggtgacggc ggagtacgtg 120
 nacgagggcg gcgccatggt gcccgtccgc gtgcacaccg tgctcatctc cacncancan 180
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 gtgatccctg agaagtacct cgacgagaag acca 274

<210> 126

<211> 260
 <212> nucleic acid
 <213> Zea mays

 <400> 126

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 tgggtgagatc accaccaagg ccaatgtcga ctacgagaag attgtcaggg agacatgccg 120
 caacattggt ttogtgtcga acgatgtcgg gcttgacgt gaccactgca aggtgcttgt 180
 gaacattgag cagcagtcctt ctgatattgc tcagggtgtg cacggccact tcaccaagcg 240
 ccccgaggag attggagctg 260

<210> 127
 <211> 516
 <212> nucleic acid
 <213> Zea mays

 <400> 127

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 attcccgggt cgaccacgc gtccggagaa gaccatcttc cacctcaacc cgtccggggcg 120
 cttegtcatc ggcgggcccc aggggtgacgc cggcctcacc ggccgcaaga tcatcatcga 180
 cacgtacggc ggctggggag cccacggcgg tggcgcttc tccggcaagg accctaccaa 240
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 gctcncctcg ccgtgncctc gtgcaggtgt cgtacgcnat cggcgtgcac ggagcccntg 360
 tatcgtattc gtaactcgta cggaacnggn acgatncnng anaaggatat actanangat 420
 agtgaaggag aantnntnct tnatgcnnnn gttgatnagg atnnaanntn nannngnna 480
 angtnnnnn nnggnnnatt nnnnantnntn nnnnta 516

<210> 128
 <211> 264
 <212> nucleic acid
 <213> Zea mays

 <400> 128

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ccgcaagatc atcattgaca cctacggtgg ctggggagcc catggtggtg gcgctttctc 180
 cggcaaggac ccaaccaagg ttgaccgcag cggagcctat gtcgcaangc aggctgccaa 240
 gagcatcgtc gccagcggcc ttgc 264

<210> 129
 <211> 270
 <212> nucleic acid
 <213> Zea mays
 <400> 129

caagaatgga acctgcccct ggctcaggcc cgatgggaag acccaggtga cagtggagta 60
 ccgcaacgag ggtggcgcca tggttcccat ccgtgtgcac acagtcctca tctctaccca 120
 gcacgacgag acagtcacca acgacgagat tgctgctgac ctgaaggagc acgtcatcaa 180
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 ccgcttcgtc atcggcggac ctcacggcga 270

<210> 130
 <211> 249
 <212> nucleic acid
 <213> Zea mays
 <400> 130

cacgtcatca agcccgtgat ccttgagaag tacctcgacg agaagaccat cttccacctc 60
 aaccgctccg ggcgcttcgt catcggcggg cccacagggtg acgccggcct caccggccgc 120
 aagatcatca tcgacacgta cggcggctgg ggagcccacg gcggtggcgc cttctccggc 180
 aaggacccca ccaaggtgga ccgcagcggc gcctacgtgg ccaggcaggc cgccaagagc 240
 atcgtggcc 249

<210> 131
 <211> 270
 <212> nucleic acid
 <213> Zea mays
 <400> 131

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 ccattctggcc gctttgttat tgggtggacct cacggcgatg ctggcctcac tggccgcaag 120

atcatcattg acacctacgg tggctgggga gcccatgggtg gtggcgcttt ctccggcaag 180
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gtcgccagcg gccttgctcg ccgcgccatc 270

<210> 132
<211> 265
<212> nucleic acid
<213> Zea mays
<400> 132

ctgcancaag accaacaatgg tcatggtctt tggtgagatc accaccaagg ccaatgttga 60
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gcttgacgcc gaccactgca aggtgctcgt gaacattgag cagcagtccc ctgatattgc 180
tcaggggtgtg canggccact tcaccaagcg ccccgaggag attggagctg gtgaccaggg 240
acacatgttc gggatatgca ccgat 265

<210> 133
<211> 284
<212> nucleic acid
<213> Zea mays
<400> 133

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tctccggcaa ggacccacc aaggtggacc ncagcggcgc ctacgtggcc aggcaggccg 120
ccaagagcat cgtngccagc ngcctcgccc gccgctgcct cgtgcagggtg tcgtacgcca 180
toggntgccg gagncctgt ccgtgttcgt caactcgtac ggcaccggca cgatccccga 240
caaggagatc ctcaagatcg tgaaggagna ttcgattcag gccg 284

<210> 134
<211> 429
<212> nucleic acid
<213> Zea mays
<400> 134

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gcaaggtgct tgtgaacatt gagcagcagt ccctgatat tgctcagggt gtgcacggcc 120

acttaccaag cgccccgagg agattggagc tggtgaccag gggcacatgt ttgggtatgc 180
gactgacgag acccctgagc tgatgccct cagccatgtn cttgccacca agcttggtgc 240
tcgtctnaca aangntcgca agaaatggaa cctggcccct ggcttaagcc cgatnggnaa 300
gaccaagtgc acaagtggaa tanccgnaac caaggggtggc nccatgggtt cccattcgtg 360
tgcacacaag tccttaattt ttacccaaca ccgaccaagg ccagttancc aacgaccagg 420
anttggcnt 429

<210> 135
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 135

atcggcgggc cccacggtga cgccggcctc accggccgca agatcatcat cgacacgtac 60
ggcggctggg gagccacgg cggtggcgcn ttctccggca aggacccac caaggtggac 120
cgcagcggcg cntacgtggc caggcaggcc gccaaagca tcgtngccag cngctcgnc 180
gccgntgcnt nggtgcaggtg tcgtacgcca tcggctgccg gagcccctgt ccgtgttngt 240
caactcgtac ggcnegcgga cgnccccga caaggagntc tc 282

<210> 136
<211> 279
<212> nucleic acid
<213> Zea mays

<400> 136

gtgatggtgt tcggcgagat caccaccaag gcgaccgtgg actacgagaa gatcgtgcgc 60
gacacctgcc gcgagatcgg gttcacctcc gacgacgtgg gcctcgacgc cgaccgctgc 120
aaggtgctgg tgaacatcga gcagcagtc cccgacatcg cgcagggcgt gcacgggcat 180
tcacgaagcg gcccaggag atcggcgcg gcgaccagg ccacatgttc gggtagcca 240
ccgacgagac ccccagtgga tgccgtnagc natgtngc 279

<210> 137
<211> 283
<212> nucleic acid
<213> Zea mays

<400> 137

ctgagaagta cctcgacgag aagaccatct tccacctcaa cccgtccggg cgcttcgtca 60
ttggcggggc ccaaggtgaa ggccgggctta acgggcggaa anntnntcat cganacgnan 120
ggcgggttggg gagccacagg cgggtggcgn ttctccggca aggacccac caaggtggac 180
cgcagcggcg cctacgtggc caggcaggcc gccaaagca tcgtggccag cggcttcgcc 240
cgcngctgcc tcgtgcaggt gtcgtacgcc atcgggtgcc gga 283

<210> 138

<211> 297

<212> nucleic acid

<213> Zea mays

<400> 138

cggaentggn gaaaggagca cgtcatcaag ccagtcattc ccgagcagta cctcgacgag 60
annntcaatc ttccacctca acccgtctgg ccgcttcgtc atcggcggac ctcacggcga 120
cgctggcctc actggccgga agatcatcat cgacacctac ggtggctggg gagccacagg 180
cgggggcgcc ttctccggca aggacccgac caaggtggac cgcagcgggg cctacgtcgc 240
gaggcaggct gccaaagca tcgtcgccgc ggcttcgcc gccgcgatt gtccagt 297

<210> 139

<211> 317

<212> nucleic acid

<213> Zea mays

<400> 139

ctccctcttg ccggtcccga ataaagagca gcagcgcaag aggtcggtag agcgagaaga 60
aggcaatggc ggccgagagc ttccctttca cctcggagtc cgtgaacgag gggcaccoga 120
caagctgtgc gaccaggtgt cggacgccgt gcttgacgca tgcctcggc aggacccoga 180
cagcaagggtg gcctgcgaga cctgcaccaa gaccaacatg gtgatgggtg tcggcgagat 240
cacgaccaag gcgaccgtgg actacgagaa gatcgtgcgc gacacctgcc gcgagatcgg 300
gttcacctcc gacgacg 317

<210> 140

<211> 277

<212> nucleic acid

<213> Zea mays

<400> 140

cgctgcctcg tgcaggtgtc gtacgccatc ggctgtggccg gagccctgt ccgtgttctg 60
caactcgtac ggcaccggca cgatccccga caaggagatc ctcaagatcg tgaaggagaa 120
cttcgacttc aggcccgga tgatcagcat caacctcgac ctgaagaagg gcggcaacag 180
gttcatcaag accgccgctt acggccactt cggccgtgac gacgccgact tcacctggga 240
ggtggtgaag cccctcaagt tcgacaaggc atcgctt 277

<210> 141

<211> 279

<212> nucleic acid

<213> Zea mays

<400> 141

ccgagcctct ctccgtgttc gtcgacacgt acggcaccgg cgcgatcccc gacaaggaga 60
tcctcaagat tgtcaaggag aacttcgatt tcaggcctgg catgatcatc atcaaccttg 120
acctcaagaa aggcggcaac gggcgctacc tcaagacggc ggccctacggc cactttggaa 180
gggacgaccc tgacttcacc tgggaggtgg tgaagccact caagtggag aaaccttctg 240
cctaaggcgg ctttttttcc agtaagaagc ttttggtgg 279

<210> 142

<211> 263

<212> nucleic acid

<213> Zea mays

<400> 142

caaggttgac cgcagcggag cctatgtcgc aaggcaggct gccaaagca tcgtcgccag 60
cngccttgct cgccgcgcca tcgtccaggt gtcttacgcc atcggcgtgc ccgagcctct 120
ctccgtgttc gtcgacacgt acggcaccgg cgcgatcccc gacaaggaga tcctcaagat 180
tgtcaaggag aacttcgatt tcaggcctgg catgatcatc atcaaccttg acctcaagaa 240
aggcggcaac gggcgctacc tca 263

<210> 143

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 143

cgatcatcaag ccagtcattcc ccgagcagta cctcgacgag aagacaatct tccacctcaa 60
 cccgtctggc cgtttctgtca tcggcggacc tcacggcgac gccggcctca ctggccggaa 120
 gatcatcatc gacacctacg gtggctgggg agcccacggc gggggcgctt tctccggcaa 180
 ggacccgacc aaggtggacc gcagcggggc ctacgtcgcg aggcaggctg ccaagagcat 240
 cgtcgccgcc ggccttcgcc cgcngcgcca tcgtccaggt ctctaag 287

<210> 144

<211> 280

<212> nucleic acid

<213> Zea mays

<400> 144

ggccacttca ccaagcgccc cgaggagatt ggagctgntg accagggaca catgttcggg 60
 tatgcgaccg atgagacccc tgagttgatg ccctcagcc atgtccttgc caccaagcta 120
 ggtgctcgtc tcaccgaggt ccgcaagaac ggaacctgcc cctggctcag gcctgatggg 180
 aagaccaggt tgacagtoga gtaccgcaat gaggggtggg ccatgggtccc catccgtgtc 240
 cacaccgtcc tcattctcac ccgcacgacg agacagtgac 280

<210> 145

<211> 251

<212> nucleic acid

<213> Zea mays

<400> 145

ttcacctccg acgacgtggg cctcgacgcc gaccgctgca aggtgctggt gaacatcgag 60
 cagcagtcac ccgacatcgc gcagggcgtg caggggcact tcacgaagcg gcccgaggag 120
 atcggcgcgg ggcaccaggg ccacatgttc gggtagccca ccgacgagac ccccgagctg 180
 atgccgctga gccacgtgct ggccaccaag ctggggcgcg gcctcaccga ggtgcgcaag 240
 aacggcactg g 251

<210> 146

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 146

atggtcccca tccgtgtcca caccgtcctc atctccaccc agcacgacga gacagtgacc 60
aatgatgaga tccgtgtctga cctgaaggag catgtcatca agcctgtcat ccctgagcag 120
taccttgacg agaagaccat ctccacctt aaccatctg gccgtttgt cattgggtgga 180
cctcacggcg atgctggcct cactggccgc aagatcatca ttgacaccta cgggtggctgg 240
ggagcccatg gtggtggcgt ttctccggca 270

<210> 147

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 147

agacctgccg caacattggt ttgtgtcaa acgatgttg gcttgacgcc gaccactgca 60
aggtgctcgt gaacattgag cagcagtcct ctgatattgc tcagggtgtg catggccact 120
tcaccaagcg ccccgaggag attggagctg gtgaccagg acacatgttc gggatatgca 180
ccgatgagac ccctgagttg atgcccctca gccatgtcct tgccaccaag ctaggtgctc 240
gtctcaccga ggtccgcaag aacggaactg cccctggctc agcctgatgg gaagaccagt 300
gacagtcgag 310

<210> 148

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 148

ccagggtctcc tacgccatcg gcgtgcccga gcccctttcg gtgttcgtgg acacgtacgg 60
caccggcgcg atccccgaca aggagatcct gaagatcgtg aaggagaact togacttcag 120
gcccggcatg atcatcatca acctcgacct caagaaaggc ggcaacgggc gctacctcaa 180
gacggcgggc tacgggcact ttgggaggga cgaccccgac ttcacctggg aggtgggtgaa 240
gcccctcaag gcggagaagc cgtcttctgc atgaggcgcc tcctctgttt ng 292

<210> 149

<211> 279
 <212> nucleic acid
 <213> Zea mays

 <400> 149

 aggtgacagt cgagtaccgc aatgaggggtg gtgccatggt ccccatgcgt gtccacaccg 60
 tcctcatact ccacccagca cgacgagaca gtgaccaatg atgagatcgc tgctgacctg 120
 aaggagcatg tcatacaaggc tgtcatcctg agcagtacct tgacgagaag accatcttcc 180
 accttaacnc atctggccgc tttgtcattg gtggacctca cggcgatgct ggccctcactg 240
 gccgcaagat catcattgac acctacggtg gctggggag 279

<210> 150
 <211> 322
 <212> nucleic acid
 <213> Zea mays

 <400> 150

 ctccctcttg ccgggtccga ataaagagca gcagcgcaag aggtcggtag agcgagaaga 60
 aggcaatggc ggccgagagc ntctttttca cctcggagtc cgtgaacgag gggcaccctg 120
 acaagctgtg cgaccaggtg tcggacgccg tgcttgacgc atgcctcgcg caggaccctg 180
 acagcaaggt ggctgcgag acctgcacca agaccaacat ggtgatgggtg ttcggcgaga 240
 tcacgaccaa ggcgaccgtg gacnacgaga agatcgtgcg cgacacctgc cgcgagatcg 300
 ggttcactcc gacgacgtgg gc 322

<210> 151
 <211> 283
 <212> nucleic acid
 <213> Zea mays

 <400> 151

 gaataaagag cagcagcgca agaggtcgggt agagcgagaa gaaggcaatg gcggccgaga 60
 gcttcctttt cacctcggag tccgtgaacg aggggcaccc cgacaagctg tgcgaccagg 120
 tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc cgacagcaag gtggcctgcg 180
 agacctgcac caagaccaac atggtgatgg tggtcggcga gatcacgacc aaggcgaccg 240
 tggactacga gaagatcgtg cgcgacacct gccgagat cgg 283

<210> 152
 <211> 316
 <212> nucleic acid
 <213> Zea mays

 <400> 152

 cgtttgccctc ttctccctct tgccggtccc gaataaagag cagcagcgca agagggtcggt 60
 agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
 aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
 cgcaggaccc cgacagcaag gtggcctgcg agacctgcac caagaccaac atggtgatgg 240
 tgttcggcga gatcacgacc aaggcgaccg tggactacga gaagatcgtg cgcgacacct 300
 gccgcgagat cggggtt 316

<210> 153
 <211> 277
 <212> nucleic acid
 <213> Zea mays

 <400> 153

 gcatggccac ttcaccaagc gccccgagga gattngagct ggtgaccagg gacacatggt 60
 ccgggtatgc gaccgatgag acccctgagt tgatgccctt cagccatgtc cttgccacca 120
 agctaggtgc tcgtctcacc gaggtccgca agaacggaac ntgcccctgg ctcagggctg 180
 atgggaagac ccaggtgaca gtcgagtacc gcaatgaggg tggtgccatg gtcccatcc 240
 gtgtccacac cgtcctcatc tccaccacgc acgacga 277

<210> 154
 <211> 272
 <212> nucleic acid
 <213> Zea mays

 <400> 154

 tgtcatcaag cctgtcatcc ctgagcagta ccttgacgag aagaccatct tccaccttaa 60
 cccatctggc cgctttgtca ttggtggacc tcacggcgat gctggcctca ctggccgcaa 120
 gatcatcatt gacacctacg gtggctgggg agcccatggt ggtggcgctt tctccggcaa 180
 ggaccaaac aaggttgacc gcagcggacc tatgtcgcaa ggcaggctgc caagagcatc 240

gtcgccagcg gccttgctcg ccgcgccatc gt

272

<210> 155
<211> 297
<212> nucleic acid
<213> Zea mays

<400> 155

ccctcttgcc ggtcccgaat aaagagcagc agcgcaagag gtcggtagag cgagaagaag 60
gcaatggcgg ccgagagctt ccttttcacc tcggagtccg tgaacgaggg gcaccccgac 120
aagctgtgcg accaggtgtc ggacnccgtg cttgacgcat gcctcgcgca ggaccccgac 180
agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtggt cggcgagatc 240
acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacctgccg cgagatc 297

<210> 156
<211> 267
<212> nucleic acid
<213> Zea mays

<400> 156

gtcggtagag cgagaagaag gcaatggcgg ccgagagctt ccttttcacn tcggagtccg 60
tgaacgaggg gcaccccgac aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat 120
gcctcgcgca ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg 180
tgatggtggt cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg 240
acacctgccg cgagatcggg ttacact 267

<210> 157
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 157

aggagattgg agctggtgac caggggcaca tgtttggtga tgcgactgac gagaccctg 60
agctgatgcc cctcagccat gtccttgcca ccaagcttgg tgctcgtctc acggagggtc 120
gcaagaatgg aacctgcccc tggctcaggc ccgatgggaa gaccaggtg acagtggagt 180
accgcaacga ggggtggcgcc atggttccca tccgtgtgca cacagtctc atctctaccc 240

agcacgacga gacagtcacc a

261

<210> 158
<211> 288
<212> nucleic acid
<213> Zea mays

<400> 158

gcccctcagc catgtccttg ccaccaagct aggtgctcgt ctcaccgagg tccgcaagaa 60
cggaacctgc ccttggtca ggctgatgg gaagaccag gtgacagtcg agtaccgcaa 120
tgaggggtgg gccatggtcc ccacccgtgt ccacaccgtc ctcactcca cccagcacga 180
cgagacagtg accaatgatg agatcgctgc tgacctgaag gagcatgtca tcaagcctgt 240
catccctgag cagtacttga cgagaagaca tttccactt aaacccat 288

CCAGGTCGG TAGAGCGAGA
AGAAGGCAAT GGCGGCCGAG AGCTTCTTT TCACNTCGGA GTCCGTGAAC GAGGGGCACC
CGACAAGCT GTGCGACCAG GTGTCGGACG CCGTGCTTGA CGCATGCCTC GCGCAGGACC
CGACAGCAA GGTGGCNTGC GAGACNTGCA CCAAGACCAA CNTGGTGATG GTGTTGCGCG
AGATCACGAC CAAGGCGACC GTGGACTACG AGAAGATCGT GCGCGACACC TGCCGCGAGA
TCGGGTTTAC T

<210> 159
<211> 311
<212> nucleic acid
<213> Zea mays

<400> 159

cttctccctc ttgccggtcc cgaataaaga gcagcagcgc aagaggtcgg tagagcgaga 60
agaaggcaat ggcggccgag agcttcttt tcacntcgga gtccgtgaac gaggggcacc 120
cgcacaagct gtgcgaccag gtgtcggacg cctgtcttga cgcattgctc ggcgaggacc 180
cgacagcaa ggtggcntgc gagacntgca ccaagaccaa cntggtgatg gtgttcggtg 240
agatcacgac caaggcgacc gtggactacg agaagatcgt ggcgacacc tgccgcgaga 300
tcgggttac t 311

<210> 160
<211> 267
<212> nucleic acid
<213> Zea mays

<400> 160

cggacgctgg ngcgggtggcg ccttctccgg caaggacccc accaagggtg accgcagcgg 60
cgctaactg ggccaggcag gccgccaaga gcacgtggc cagcggcctc gccgcccgt 120
gcctcgtgca ggtgtcgtac gccatcggct gccggagccc ctgtccgtgt tcgtcaactc 180

gtacggcacc ggcacgatcc ccgacaagga gatcctcaag atcgtgaagg agaacttcga 240
cttcaggccc gggatgatca gcatcaa 267

<210> 161
<211> 284
<212> nucleic acid
<213> Zea mays

<400> 161

agctgtcgac acattcctct tcacctcgga gtctgtgaac gagggacacc ctgacaagct 60
ctgtgaccag gtctcagatg ccgttcttga cgcttgctt gctgaggacc ctgacagcaa 120
ggttgcttgt gagacctgca ccaagaccaa cattcancat ggtctttggt gagatcacca 180
ccaaggccaa tgttgactac gagaagattg tcaggggagac ctgccgcaac attggttttg 240
tgtcaaacga tgttggttg acgccgacca ctgcaagggtg ctcg 284

<210> 162
<211> 237
<212> nucleic acid
<213> Zea mays

<400> 162

ccatcttcca cctcaaccgg tccgggctgt tcgtcatcgg cgggccccac ggtgacgccg 60
gcctcaccgg ccgcaagatc atcatcgaca cgtacggcgg ctggggagcc cacggcggtg 120
gcgccttctc cggcaaggac cccaccaagg tggaccgcag cggcgcttac gtggccaggc 180
aggccgcca gagcatcgtg gccagcggcc tcgcccgcg ctgcctcttg naggttt 237

<210> 163
<211> 236
<212> nucleic acid
<213> Zea mays

<400> 163

acctgcgcct ggctgaggcc cgacggcaag acccaggtga cggaggagta cgtgaacgag 60
ggcggcgcca tgggtgccgt ccgcgtgcac accgtgctca tctccacca gcacgacgag 120
accgtcacca acgacgagat cgccgccgac ctcaaggagc acgtcatcaa gcccgatgc 180
cctgagaagt acctcgacga gaagaccatc ttccacctca acccgccgg gcgctt 236

<210> 164
 <211> 272
 <212> nucleic acid
 <213> Zea mays

 <400> 164

 ggtggacctc acggcgatgc tggcctcact ggccgcaaga tcatcattga cacctacggt 60
 ggetggggag cccatggtgg tggcgctttc tccggcaagg acccaaccaa ggttgaccgc 120
 agcggagcct atgtcgcaag gcaggctgcc aagagcatcg tcgccagcgg ccttgctcgc 180
 cgcgccatcg tccaggtgtc ttacgccatc ggntggcccg agcctctctc cgtgttcgtc 240
 gacacgtacg gcaccggcgc gatccccgac aa 272

<210> 165
 <211> 258
 <212> nucleic acid
 <213> Zea mays

 <400> 165

 ccaagctagg tgctcgtctc accgaggncg gcaagaacgg aacctgcccc tggctcaggc 60
 ctgatgggaa gaccaggtg acagtcgagt accgcaatga ggggtgtgcc atggtcccca 120
 tccgtgtcca caccgtcctc atctccaccc agcacgacga gacagtgacc aatgatgaga 180
 tcgtgctga cctgaaggag catgtcatca agcctgtcat ccctgagcag taccttgacg 240
 agaagaccat cttccacc 258

<210> 166
 <211> 298
 <212> nucleic acid
 <213> Zea mays

 <400> 166

 tctccctctt gccggtcccg aataaagagc agcagcgcaa gaggtcggta gagcgagaag 60
 aaggcaatgg cggccgagag cttccttttc acctcgaggt ccgtgaacga ggggcacccc 120
 gacaagctgt gcgaccaggt gtcggacgcc gtgcttgacg catgcntcgc gcaggacccc 180
 gacagcaagg tggcctgcga gacctgcacc aagaccaaca tggatgatgt gttcggcgag 240
 atcacgacca aggcgaccgt ggactacgag aagatcgtgc gcgacacctg ccgcgaga 298

<210> 167
 <211> 298
 <212> nucleic acid
 <213> Zea mays

 <400> 167

 cttctccctc ttgccgtcc cgaataaaga gcagcagcgc aagaggtcgg tagagcgaga 60
 agaaggcaat ggcggccgag agcttccttt tcacctcgga gtccgtgaac gaggggcacc 120
 ccgacaagct gtgcgaccag gtgtcggacg ccgtgcttga cgcattgcctc gcgcaggacc 180
 ccgacagcaa ggtggcctgc gagacctgca ccaagaccaa catggtgatg gtgttcggcg 240
 agatcacgac caaggcgacc gtggactacg agaagatcnt gcgcgacacc tgccgcga 298

<210> 168
 <211> 265
 <212> nucleic acid
 <213> Zea mays

 <400> 168

 ccgagggtccg caagaacgga acctgcccct ggctcaggcc tgatgggaag acccaggtga 60
 cagtgcagta ccgcaatgag ggtgggtgcca tggccccat ccgtgtccac accgtcctca 120
 tctccacca gcacgacgag acagtgacca atgatgagat cgctgctgac ctgaaggagc 180
 atgtcatcaa gcctgtcatc cctgagcagt accttgacga gaagaccatc ttccacctta 240
 acccatctgg ccgtttgtca ttggt 265

<210> 169
 <211> 251
 <212> nucleic acid
 <213> Zea mays

 <400> 169

 cccgacaagc tgtgcgacca ggtgtcggac gccgtgcttg acgcatgcct cgcgcaggac 60
 cccgacagca aggtggcctg cgagacctgc accaagacca acatggtgat ggtgttcggc 120
 gagatcacga ccaaggcgac cgtggactac gagaagatcg tgcgcgacac ctgccgcgag 180
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 catcgagcag c 251

<210>	170
<211>	305
<212>	nucleic acid
<213>	Zea mays

ctottgccgg	tcccgaataa	agagcagcag	cgcaagaggt	cggtagagcg	agaagaaggc	60
aatggcggcc	gagancttcc	ttttnanctc	ggaatncgtg	aacgaggggc	ancccgacaa	120
gctgtgcgac	caggtgtcgg	acgccgtgct	tgacgnatgc	ctcgcgcagg	accccgacag	180
caaggtggcc	tgcgagacct	gcaccaagac	caacatggtg	atggtgttcg	gcgagatcac	240
gaccaaggcg	accgtggact	acgagaagat	cgtgcgcnac	acctgcccg	agatcggggt	300
cactc						305

<210>	171
<211>	267
<212>	nucleic acid
<213>	Zea mays

cntacggttg	ctggggagcc	cacggcgggg	gcgctttctc	cggcaaggac	ccgaccaagg	60
tggaccgcag	cggggcctac	gtncngnggc	aggctgccaa	gagcatcgtc	gcgcgcggcc	120
tcgcccgcgc	tgccatcgtc	cagggtctcct	acnccatcgg	cgtgccegan	cccctgtcgg	180
tgttcgtgga	cacgtacggc	accggcgcgga	tccccgacaa	ggagatcctg	aagatcgtga	240
aggagaactt	cgacttcagg	cccggca				267

<210>	172
<211>	250
<212>	nucleic acid
<213>	Zea mays

ccaatgatga gatcgctgct gacctgaagg agcatgtcat caagcctgtc atccctgagc	60
agtaccttga cgagaagacc atcttcacacc ttaacccatc tggcgcgttt gtcattgggtg	120
gacctcacgg cgatgctggc ctcaactggcc gcaagatcat cattgacacc tacggtgggt	180
ggggagccca tgggtgggtggc gctttctccg gcaaggaccc aaccaagggt gaccgcagcg	240

ganctatgtc

250

<210> 173
<211> 304
<212> nucleic acid
<213> Zea mays

<400> 173

ctcctccctc ctgccgggtc cttataaaag agcagcagcg caanganctg gtagancgag 60
cgagaagaag gcaatggcgg cggagagctt cctgttcacc tcggagtccg tgaacgaggg 120
gcacccagac aagctgtgcg accaggtgtc ggacgcggtg ctggacgcct gcctggcgca 180
ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag acgaacatgg tgatggtgtt 240
cggcgagatc accaccaagg cgagcgtgga ctacgagaag atcgtgcgcg acacctgccg 300
cgag 304

<210> 174
<211> 328
<212> nucleic acid
<213> Zea mays

<400> 174

catccgtttg cctcttctcc ctcttgccgg tcccgaataa agagcagcag cgcaagaggt 60
cggtagagcg agaagaaggc aatggcgggc gagagcttcc ttttcacctc ggagtccgtg 120
aacgangggc accccgacaa gctgtgcgaa ccaggtgtcg gacgccgtgc ttgacgcatg 180
cctgcgcgag gaccccgaca gcaangtggc ctgcgagacc tgcaccaaga ccaacatggt 240
gatggtgttc ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga 300
cacctgccgc gagatcgggt tcactccg 328

<210> 175
<211> 297
<212> nucleic acid
<213> Zea mays

<400> 175

tctccctctt gccggtcccg aataaagagc agcagcgcaa gaggtcggta gagcgagaag 60
aaggcaatgg cggccgagag ctctcttttc acctcgaggt ccgtgaacga ggggcacccc 120

gaacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc 180
cgacagcaag gtggcctgcg agacctgcac caagaccaac atggatgatgg tgttcggcga 240
gatcacgacc aaggcgaccg tggactacga gaagatcgtg cgcgacacct gccgcga 297

<210> 176
<211> 275
<212> nucleic acid
<213> Zea mays
<400> 176

agatgccgtt cttgacgctt gccttgctga ggaccctgac agcaagggtt cttgtgagac 60
ctgcaccaag accaacaatgg tcatggtctt tggtgagatc accaccaagg ccaatgttga 120
ctacgagaag attgtcaggg agacctgccg caacattggt tttgtgtcaa acgatgttgg 180
gcttgacgcc gaccactgca aggtgctcgt gaacattgag cagcagtccc ctgatatgct 240
caggggtgtgc atggccattc accaagcgcc ccgag 275

<210> 177
<211> 534
<212> nucleic acid
<213> Zea mays
<400> 177

gaggnagctt tnannggggn ttaagggnat ttttaaacc tnnnttgnat tcccgggtcg 60
acccacgcgt ccgcttgccg gtcccgaata aagagcagcn ncgcaagagg tcggtagagc 120
gagaagaagg caatggcggc cgagagcttc cttttcacct cggagtccgt gaacgagggg 180
caccocgaca agctgtgcga ccaggtgtcg gacgccgtgc ttgacgcatg cctcgcgcag 240
gaccccgaca gcaagggtggc ctgcgagacc tgcaccaaga ccaacatggt gatggtgttc 300
ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga caccttnccg 360
cgagatcggg ttcacctttc gacgacntgg gccttgacgc ccaccgggtt caaggtgctt 420
gtgnacattg agcaagaatt ccccgaaatt gngcaaggcg ttcaccggca ctttacgaac 480
cggcccnagg aagatcggnc cggccnacca nggncaattt tttgggtccc cccc 534

<210> 178
<211> 248
<212> nucleic acid

<213> Zea mays

<400> 178

gatgcccctc agccatgtcc ttgccaccaa gcttggtgct cgtctcacgg aggttcgcaa 60
gaatggaacc tgcccctggc tcaggcccga tgggaagacc caggtgacag tggagtaccg 120
caacgagggg ggcccatgg ttcccatccg tgtgcacaca gtccctcatct ctaccagca 180
cgacgagaca gtcaccaacg acgagattgc tgctgacctg aaggagcacg tcatcaagcc 240
agtcatcc 248

<210> 179

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 179

ctcctccctc ctgccgggtc cttataaaag agcagcagcg caagagggttg gtagagcgag 60
cgagaagaag gcaatggcgg cggagagcct cctgttcacc tcggagtccg tgaacgaggg 120
gnaccagac aagctgtgcg accaggtgtc ggacgcggtg ctggacgcct gcctggcgca 180
ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag acgaacatgg tgatggtgtt 240
cggcgagatc accaccaagg cgagcgtgga ctacgagaag atcgtgcgcg acacctgccg 300
cg 302

<210> 180

<211> 281

<212> nucleic acid

<213> Zea mays

<400> 180

angtctcta cgccatcggc gtggcccgan cccctgtcgg tgttcgtgga cacgtacggc 60
accggcgcgga tccccgacaa ggagatcctg aagatcgtga aggagaactt cgacttcagg 120
cccggcatga tcatcatcaa cctcgacctc aagaaaggcg gcaacgggcy ctacctcaag 180
acggcggcct acgggcactt tgggaggagc gaccccgact tcacctggga aggtggtgaa 240
gcccctcaag gcggagaagc cgtcttctgc atgaggcgcn t 281

<210> 181

<211> 269
 <212> nucleic acid
 <213> Zea mays

 <400> 181

 gacacattcc tcttcacctc ggagtctgtg aacgagggac accctgnana agctctgtga 60
 ccaggtctca gatgccgttc ttgacgcttg ccttgctgag gaccctgaca gcaaggttgc 120
 ttgtgagacc tgcaccaaga ccaacatggt catggtcttt ggtgagatca ccaccaaggc 180
 caatggtgac gccgagaaga ttgtcagggg gacctgccgc aacattgggt ttgtgtcaaa 240
 cgatggtggg cttgacgcng accatgcaa 269

<210> 182
 <211> 286
 <212> nucleic acid
 <213> Zea mays

 <400> 182

 ccctcttgcc ggtcccgaat aaagagcagc agcgcaagag gtcggtagag cgagaagaag 60
 gcaatggcgg ccgagagctt ccttttcacc tcggaanccg tgaacgaggg gcaccccgac 120
 aagotgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180
 agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtggt cggcgagatc 240
 acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacct 286

<210> 183
 <211> 240
 <212> nucleic acid
 <213> Zea mays

 <400> 183

 gcgcgacacc tgccgcgaga tcgggttcac ctccgacgac gtgggcctcg acgccgaccg 60
 ctgcaaggtg ctggtgaaca tcgagcagca gtcccccgac atcgcgcagg gcgtgcacgg 120
 gcaacttcacg aagcggccccg aggagatcgg cgcgggcgac cagggccaca tggtcgggta 180
 cgnacccgac gagacccccg agctgatgcc gctgagccac gtggtggcca ccaagctggg 240

<210> 184
 <211> 250
 <212> nucleic acid

<213> Zea mays

<400> 184

cctgagcagt accttgacga gaagaccatc ttccacctta acccatctgg ccgctttgtc 60
attggtggac ctacaggcga tgctggcctc actggccgca agatcatcat tgacacctac 120
ngtggctggg gagcccatgg tgggtggcgt ttctccggca aggacccaac caagggtgac 180
cgcagcggag cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct 240
cgccgcgcca 250

<210> 185

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 185

gcctcttctc ctccctcctg ccgggtcctt aataaagagc agcagcgcaa gaggttggtg 60
gagcgagcga gaagaaggca atggcgggcg agagcttcct gttcacctcg gagtccgtga 120
acgaggggca cccagacaag ctgtgcgacc aggtgtcgga cgcggtgctg gacgcctgcc 180
tggcgcagga ccccgacagc aagggtggcct gcgagacctg caccaagacg aacatgggtga 240
tggtgttcgg cgagatcacc accaaggcga gcgtggacta cgagaagatc gtgcgcgaca 300
cctgc 305

<210> 186

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 186

cgacagcaga ggnggcnngc gagacctgca ccaagaccaa catggtgatg ttgttcggcg 60
agatcacgac caaggcganc gtggactacg agaagatcgt gcgcgacacc tgatcgcgag 120
atcgggttca cctcccgacg acgtgggcct cgacgccgac cgctgcnagg ngctgggtgaa 180
natcgagcan cagtcncccc acatcgcgca ngcntgcacg ggcacttcac naagcgnccc 240
gangagatcg ncgcggcncta ccatnggcac atgttcgggt acnncaccna nnagacnnnc 300
gagct 305

<210> 187
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 187

 ccttgccacc aagcttggtg ctctgtctcac ggagggttcgc aagantggaa cctgcccctg 60
 gotcaggccc gntgggaaga ccaggtgac agtggagtac cgcaacgagg gtggcgccat 120
 ggttcccata cgtgtgcaca cagtctcat ctctaccag cagcagaga cagtcancaa 180
 cgaagagatt gctgctgacc tgaaggagca cgtcatcaag ccagtcattc ccgagnagnn 240
 acctcgacga gaagacaatc ttccacacac ttna 274

<210> 188
 <211> 232
 <212> nucleic acid
 <213> Zea mays

 <400> 188

 agcagtcatt caagccagtc atccccgagc agtacctcga cgagaagaca atcttccacc 60
 tcaaccgctc tggcgcgttc gtcctcggcg gacctcacgg cgacgctggc ctactggcc 120
 ggaagatcat catcgacacc tacggtggct ggggagccca cggcgggggc gccttctccg 180
 gcaaggaccc gaccaaggtg gaccgcagcg gggcctacgt cgcgaggcag gc 232

<210> 189
 <211> 243
 <212> nucleic acid
 <213> Zea mays

 <400> 189

 gccctcagc catgtccttg ccaccaagct tgggtgctgt ctacagagg ttgcgaagaa 60
 tggaaacctgc ccctggctca ggccgatgg gaagaccag gtgacagtgg agtaccgcaa 120
 cgagggtggc gccatggttc ccatccgtgt gcacacagtc ctcatctcta ccagcacga 180
 cgagacagtc accaacgacg agattgctgc tgacctgaag gagcacgtca tcaagccagt 240
 cat 243

<210> 190

<211> 290
 <212> nucleic acid
 <213> Zea mays

 <400> 190

 ccctcttgcc ggtcccgaat aaagancagc agcgcaagag gtcggtagag cnagaagaan 60
 gcaatggcgg ccgagagctt ccttttcacc tcggagtccg tgaacgaggg gcaccccgac 120
 aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180
 anc naggtgg cctgcgagac ctgcaccaag acca acatgg tgatgggtgtt cggcnagatc 240
 acgaccaagg cgaccgtgga ctacnagaag atcgtgcgcg acacctgccg 290

<210> 191
 <211> 266
 <212> nucleic acid
 <213> Zea mays

 <400> 191

 gncacattcc ncttcacctc ggngtctgtg ancgagggac accctgacna gctctgtgac 60
 caggtctcag atgccgttct tgacgcttgc cttgctgagg ancctgacag caaggttgct 120
 tgtgagacct gcaccaagac caacatggtc atggtctttg gtgagatcac caccaaggcc 180
 aatgttgact acgagangat tgtcagggag acctgccgca acattggttt tgtgtcaaac 240
 gatgttgggc tgacgccgac cactgc 266

<210> 192
 <211> 276
 <212> nucleic acid
 <213> Zea mays

 <400> 192

 gtgccatggt ccccatccgt gtccacaccg tcctcatctc caccagcac gacgagacag 60
 tgaccaatga tgagatcgct gctganctga aggagcatgt catcaagcct gtcacccctg 120
 agcagtacct tgacgagaag accatcttcc acttaaccba tctgcccgtt ttgtcattgg 180
 tggacctcac ggcgatgctg gcctcactgg ccgcaagatc atcattgaca ctacggtggc 240
 tggggagccn atgntgtggc gtttncnggg aaggcc 276

<210> 193

<211> 292
 <212> nucleic acid
 <213> Zea mays

 <400> 193

 anggtggctg gggagcccat ggtggtggcg ctttctccgg caaggaccca accaaggttg 60
 acngcagcgg agcctatgtn cgcaaggcag gctgccaaaga gcatcgtcgc cagcggcctt 120
 gctcgccgcg ccatcggtcca ggtgtcttac gccatcgggc tgcccagacc tctctccgtg 180
 ttctcgaca cgtacggcac cggcgcgatc cccgacaagg agatcctcaa gattgtcaag 240
 gagaattcga tttcaggcct ggcgatgatc catcaacctt gactcaagaa gg 292

<210> 194
 <211> 226
 <212> nucleic acid
 <213> Zea mays

 <400> 194

 ctacgagaag atcgtgctcg acacctgccg cgagatcggg ttcacctccg acgacgtggg 60
 cctcgacgcc gaccgctgca aggtgctggt gaacatcgag cagcagtcgc ccgacatcgc 120
 gcagggcgctg caggggcact tcacgaagcg gcccagaggag atcggcgcgcg gcgaccaggg 180
 ccacatgttc gggtagccca ccgacgagac ncccagagctg atgcng 226

<210> 195
 <211> 289
 <212> nucleic acid
 <213> Zea mays

 <400> 195

 cttcnccctc ttgccgtcc cgaataaaga gcagcagcgc aagaggncgg tagagcgaga 60
 agaaggcaat ggcggccgag agcttccttt tcacctcgga gtccgtgaac gaggggcacc 120
 ccgacaagct gtgcgaccag gtgtcgagcg ccgtgcttga cgcagtgcctc gngcaggacc 180
 ccgacagcaa ggtggcctgc gagacctgca ccaagaccaa catggtgatt gtgttcggcg 240
 agatcacgac canggcgacc gtggactacg agaagatcgt gcgcnacac 289

<210> 196
 <211> 300
 <212> nucleic acid

<213> Zea mays

<400> 196

cgctctttct cctccctcct gccgggtcct taataaagag cagcagcgca agnggttggt 60
agagcgagcg agaagaaggc aatggcgggc gagagcttcc tgttcacctc ggagtcctgt 120
aacgaggggc acccagacaa gctgtgcgac cagggtgcgg acgcggtgct ggacgcctgc 180
ctggcgagc accccgacag caaggtggcc tgcgagacct gcaccaagac gaacatggtg 240
atggtgttcg gcgagatcac caccaaggcg agcgtggact acgagaagat cgtgcgcgac 300

<210> 197

<211> 284

<212> nucleic acid

<213> Zea mays

<400> 197

ccctcttgnc ggtcccgaat aaagagcagc agcgcaagag gtcgntagag cgagaagaag 60
gcaatggcgg ccgagagctt ccttttcacc tcggagtccg tgaacgaggg gcaccccgac 120
aagctgtgag accaggtgtc ggacgccgtn cttgangcat gcctcgcgca ggaccccgac 180
agcaaggtgg cctgcgagac ctgnaccaag acnaacatgg tgatggtggt cggcgagatc 240
acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acac 284

<210> 198

<211> 282

<212> nucleic acid

<213> Zea mays

<400> 198

gtcccgaata aagagcagca gcgcaagang tcggtagagc ganaanaang caatggcggc 60
naagagcttc cttttcacct cggagtccgt gaacgagggg cancccgaac aagctgtncg 120
accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac agcaaggtgg 180
cctgcgagac ctgcaccaag accaacaatng tgatggtggt cggcgagatc acgaccaang 240
cgaccgtgga ctacgagaag atcgtgcgcg acacctgccg cg 282

<210> 199

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 199

acgccatcgg ntgccgganc cctgtccgt gttcgtcaac tcgtacggca ccggcacgat 60
ccccgacaag gagatcctca agatcgtgaa ggagaacttc gacttcaggc ccgggatgat 120
cagcatcaac ctgcacctga agaagggcgg caacagggttc atcaagaccg nccctacgg 180
ccacttcggc cgtgacgacg ccgacttcac ctgggagggtg gtgaagcccc tcaagttcga 240
caaggcatcg gcttaagggtt gggantgtca tgtggacatg angatacntc ct 292

<210> 200

<211> 291

<212> nucleic acid

<213> Zea mays

<400> 200

cgtttgcttc ttctccctct tgccggtccc gaataaagag cagcagcgca agaggtcggt 60
agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
angggcaccg cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
cgcaggaccg cgacagcaag gtggcctgcg agacctgcac caagaccaac atggtgatgg 240
tgttcggcga gatcacgacc aaggcgaccg tggactacga gaagatcgtg g 291

<210> 201

<211> 337

<212> nucleic acid

<213> Zea mays

<400> 201

gtnttacgcn atcngcaggc ccnagnntct ctncgtgttc gtcgacacgt anggcancgg 60
cgngatnnn ganaaggaga tcctcaagat tgtnaaggng aactnngatt tcaggcctgg 120
catgatcatc atcaaccttg acctcaagan aggcggnaac gggcgctacc tcaagacggc 180
ggattanggc cactttggaa gggangaccg tgacttcacc tgggatgtgg tnaagccact 240
caantcggag aaacctnctg cctaaggcgg nttntttttc agtaagaagc ttttggtggt 300
ctgctgtgct taatcatgcn ttatatggct tctacac 337

<210> 202

<211> 279
 <212> nucleic acid
 <213> Zea mays

 <400> 202

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 agaaggcaat ggcggccgag agcttccttt tcacctcgga gtccgtgaac gaggggcacc 120
 ccgacaagct gtgcgaccag gtgtcggacg ccgtgcttga cgcattgcctc gcgcaggacc 180
 ccgacagcaa ggtggcctgc gagacctgca ccaagacca catggtgatg gtgttcggcg 240
 agatcacgac caaggcgacc gtggactacg agaagatcg 279

<210> 203
 <211> 443
 <212> nucleic acid
 <213> Zea mays

 <400> 203

 aaantcacgg gtgcacncac gcgtcnnnac ngactnntac taacgcgtgg gcggacgcgt 60
 ggggganggt gtggacacgg ntggggacca tggcagcacc ctcatcgcg taatcgactg 120
 tcacccctga gcagtacctt gacgagaaga ccattctcca ccttaacca tctggccgct 180
 ttgtcattgg tggacctcac ggcgatgctg gcctcactgg ccgcaagatc atcattgaca 240
 cctacgggtg ctggggagcc catggtggtg gcgctttctc cggcaaggac ccaaccaagg 300
 ttgaccgcag cggagcctat gtgcgaaggc angctgcnaa gagcatcgtc gccagcgggc 360
 cttgctcgnc cnggccatcg tccaaggtgt ncttaagcca atcggcntgc ccgancctnt 420
 ctccgntttt cgtcnaaang tta 443

<210> 204
 <211> 290
 <212> nucleic acid
 <213> Zea mays

 <400> 204

 tgctctttct cctctttgcc ggtcccgaat aaagagcagc agcgcaagag gtcggtagag 60
 cgagaagaag gcaatggcgg ccgagagctt ctttttcacc tcggagtccg tgaacgangg 120
 gcaccccgac aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca 180

ggancccgac agcaaggtgg cctgcgagac ctgcaccaag accaacatgg tgatggtgtt 240
 cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgogcg 290

<210> 205
 <211> 304
 <212> nucleic acid
 <213> Zea mays
 <400> 205

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 agagcgagcg agaaagaagg catggcgggc gagagcttcc tgttcacntc ggagtccgtg 120
 aacgaggggc acccagacaa gctgtgagac caggtgtcgg acgcggtgct ggacnccctg 180
 ctggcgagcagg accccgacag caaggtggcc tgcgagaccn gcaccaagac gaacatggtg 240
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 acct 304

<210> 206
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 206

ctcttctcct cctcctgcc gtttccttaa taaagagcag cagcgcaaga ggttggtaga 60
 gcgagcgaga agaaggcaat ggcgcgag agcttctgt tcacctcgga gtccgtgaac 120
 gaggggcacc cagacaagct gtgcgaccag gtgtcggacg cgggtgctgga cgcctgcctg 180
 gcgcaggacc ccgacagcaa ggtggcctgc gagacctgca ccaagacgaa catggtgatg 240
 gtgttcggcg agatcaccac caaggcgagc gtnagctacg agaagatcgt 290

<210> 207
 <211> 247
 <212> nucleic acid
 <213> Zea mays
 <400> 207

gatcaccacc aaggccaatg ttgactacga gaagattgtc agggagacct gccgcaacat 60
 tggttttgtg tcaaacgatg ttgggcttga cgccgaccac tgcaagggtgc togtgaacat 120

tgagcagcag tcccctgata ttgctcaggg tgtgcatggc cacttcacca agcgccccga 180
ggagattgga gctggtgacc agngacacat gttcgggtat gcgaccgatg agaccctgag 240
ttgatgc 247

<210> 208
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 208

cgtttgccctc ttctccctct tgccgggtccc gaataaagag cagcagcgca agaggtcggt 60
agagcgagaa gaaggcaatg gcgggccgaga gcttcttttt cacctcggag tccgtgaacg 120
aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
cgcaggaccc cganagcaag gtggcctgcg agacctgcac caagaccaac atgggtgatgg 240
tgttcggcga gatcacgacc aaggcgaccg tggactacga gaagatcgtg c 291

<210> 209
<211> 428
<212> nucleic acid
<213> Zea mays

<400> 209

cggacgcttg ggcttttcgcg tggggtcgac acgtacgggn ccgggggggat cncgcacaag 60
gagatcctca agattgtcaa ggagaacttt tatttcaggc ctggcatgat catcatcaac 120
cttgacctca agaaaggcgg caacggggcgc tacctcaaga cggcgggcta cggccacttt 180
ggaagggacg accctgactt cacctgggag gtgggtgaagc cactcaagtc ggagaaacct 240
tctgcctaag gcggcctttt ttttcagtaa gaagcttttg gtggtctngc tgtgcttaat 300
catgctttta tatggcttct acatgttgga ggntctttct tgatctgcac cgggcttata 360
gnttggtgtg nactgcccta ataagtgtg cttatgagga ctggttctgg ttttgctgct 420
tatgtngt 428

<210> 210
<211> 295
<212> nucleic acid
<213> Zea mays

<400> 210

ccttaatnaa gngcagcagc gcaaggtgag cgcagcagctt gccccaggtt ggtagagcga 60

gcgagaagaa ggcaatggcg gcggagagct tctgttcac ctggaggtcc gtgaacgagg 120

ggcaccaga caagctgtgc gaccaggtgt cggacgcggt gctggacgcc tgctggcg 180

aggacccga cagcaaggtg gcctgcgaga cctgcaccaa gacgaacatg gtgatggtgt 240

tcgngagat caccaccaag gcgagcgtgg actacgagan gntcgtgcnc gacac 295

<210> 211

<211> 257

<212> nucleic acid

<213> Zea mays

<400> 211

ggccacttca ncaagcgtcc cgaggagatt ggagctggtg accagggacn cgtgttgcg 60

gtatgcgacc gatgagaccc ctgagttgnt gcccctcagc catgtccttg ccaccaagct 120

aggtgctcgt ctactgagg tccgcaagaa cggaacctgc ccctggctca ggcctgatgg 180

gaagaccnn gtgacagtcg agtaccgcaa tgagggtggt gcnatggtcc cngnnngtgt 240

ccanaccgtc ctcat 257

<210> 212

<211> 288

<212> nucleic acid

<213> Zea mays

<400> 212

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gaagatggcc ggactcgaca ccttcctctt cacctcgag tccgtgaacg agggacaccc 120

tgacaagctc tgcgaccagg tctcagatgc tggtctggac gcttgccttg ctgaggaccc 180

tgacagcaag gttgcttgcg agacctgcac caagaccaac atggtcatgg tctttggtga 240

gatcaccacc aaggcaatgt cgactagaga agattgtcag ggagacat 288

<210> 213

<211> 467

<212> nucleic acid

<213> Zea mays

<400> 213

ttnnnnnttcc ttnggccccn aaaggtaaaa ggactcccgg gtcgaccac gcgtcagcca 60

cgcgctccgcc cacgcgtccg cccacgcgtc cgcacacgog tccgcccacg cgtccgcccc 120

ctcctgccgg gtccttaata aagagcagca gcgcaagggtg agccgccagc ttgccccggg 180

tggtagagcg agcganaaga angcaatggc ggcgganagc ttctgttca cctcgggtcc 240

gtgaacgagg ggcaccaga caagctgtgc gaccagggtg cggacgcggg gctggcgcc 300

gcctggcgca ggaccccgac agcaagggtg cctgcgagac ctgcaccaag acgacatgg 360

gatgggtgtc ggcgagatna ccaccaaggc gagcgtggac tacgaaaaag atntgcgcga 420

aaactggccg ccaagatcgg gttcacctcc gacgacgtgg ggctcga 467

214
287
nucleic acid
Zea mays

<400> 214

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gtagagcgag aagaaggcaa tggcgggccga gagcttccct ttcacctcgg agtccgtgaa 120

cgagggggcac cccgacaagc tgtgcgacca ggtgtcggac gccgtgcttg acgcatgcct 180

cgcgaggagc cccgacagca aggtggcctg cgagacctgc accaagacca acatgggtgat 240

ggtgttcggc gagatcacga ccaaggcgat cgtggactac gagaaga 287

<210> 215
<211> 294
<212> nucleic acid
<213> Zea mays

<400> 215

ctcttctcct cctcctgnn gggtccttaa taaagagcag cagcgcaaga ngttggtaga 60

gcgancgana agaaggcaat ggcggcgag agcttctgt tcanctcgga gtccgtgaac 120

gagggggcacc cagacaagct gtgcgaccag gtgtcggacg cgggtgctgga cgctgcctg 180

gcgcaggacc ccgacagcaa ggtggcctgc gagacctgca ccaagacgaa catgggtgat 240

gtgttcggcg agatcaccac caaggcgagc gtggactacn agaagatcgt gcgc 294

<210> 216
 <211> 228
 <212> nucleic acid
 <213> Zea mays

 <400> 216

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 tgagggtggt gccatggtec ccatcgtgt ccacaccgtc ctcatctcca cccagcacga 120
 cgagacagtg accaatgatg agatcgctgc tgacctgaag gagcatgtca tcaagcctgt 180
 catccctgag cagtaccttg acgagaagac catcttccac cttaaccc 228

<210> 217
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 217

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 gggatatgca ctgacgagac ccctgagctg atgcccctca gccatgtcct tgccaccaag 120
 cttggtgctc gtccacggag gtctgcaaga atggaacctg cccctggctc aggcccgatg 180
 ggaagacca ggtgacagtg gactaccgca agaggggtggc gccatgggtc ccatccgtgt 240
 gcacacagtc ctcatctcta cccagcag 268

<210> 218
 <211> 263
 <212> nucleic acid
 <213> Zea mays

 <400> 218

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 catggtnatg gtnttcggcg agatcacgac caaggctacc gtgggactac gagaagatcg 120
 tgcgctacac ctgncgagag atcnggttca cctccgacga cgtgggcctc gacgcctacc 180
 gctncaaggt gctggtgaac atctagcagc agtccccga catnncgag ggogtgcacg 240
 ggcattnng aagcggcccg agg 263

<210> 219

<211> 256
 <212> nucleic acid
 <213> Zea mays

 <400> 219

 gagccacgg cgggggcgcc ttctccggca aggacccgac caaggtggac cgcagcgggg 60
 cctacgtcgc gaggcaggct gccaaagagca tcgtcgccgc cggcctcgcc cgccgcgcca 120
 ttgtccaggt ctctacgcc atcgggtgcc cganccctt tcggtgttcg tggacacgta 180
 cggcaccggc gcgatccccg acaaggagat cctgaagatc gtgaaggaga attcgacttc 240
 aggcccgga tgatca 256

<210> 220
 <211> 334
 <212> nucleic acid
 <213> Zea mays

 <400> 220

 annccggag cccctgtccg tgttcgtcaa ctctacggc accggcacga tccccgacaa 60
 ggagatcctc aagatcgtga aggagaactt cgacttcagg cccgggatga tcagcatcaa 120
 cctcgacctg aagaagggcg gcaacagggt catcaagacc gccgcctacg gccattcggc 180
 cgtgacgacg ccgacttcac ctgggaggtg gtgaagcccc tcaagttcga caaggcatcg 240
 gcttaagggt nggatgtcac tgtggacatg aggactactt cctctggctc tgctgttacc 300
 tgcaagcatt gtgctgtgga tgtgtgtgtt tgat 334

<210> 221
 <211> 255
 <212> nucleic acid
 <213> Zea mays

 <400> 221

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 cccnncanc nnggtggcct gcgagacctg caccaagacc aacatngtga tgggtgttcgg 120
 cgagatcacg ancaaggcga ccgtggacta cgagnngatc gtncgcgaca cctgccgcga 180
 gatcgggttc aatccganga ngtgggcctc nacgccgnnc gctgcnagggt gctgggtgaac 240
 ntcgagcagc agtcc 255

<210> 222
 <211> 222
 <212> nucleic acid
 <213> Zea mays
 <400> 222
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 tcatcaagcc tgtcatccct gagcagtacc ttgacgagaa gaccatcttc caccttaacc 120
 catctggccg ctttgtcatt ggtggacctc acggcgatgc tggcctcact ggccgcaaga 180
 tcatcattga cacctacggt ggctggggag cccatggtgg tg 222

<210> 223
 <211> 269
 <212> nucleic acid
 <213> Zea mays
 <400> 223
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 aggcaatggc ggccgagagc ttctttttca cctcggagtc cgtgaacgag gggcaccgcc 120
 acaagctgtg cgaccaggtg tcggacgccg tgcttgacgc atgcctcgcg caggaccgcc 180
 acagcaaggt ggctgcgag acctgcacca agaccaacat ggtgatggtg ttcggcgaga 240
 tcacgaccaa ggcgaccgtg gactacgag 269

<210> 224
 <211> 311
 <212> nucleic acid
 <213> Zea mays
 <400> 224
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 ggttgggaga gcgagcgaga agaaggcaat ggcgggcgag agcttctgt tcacctctga 120
 gtccgtgaac gangggcanc cagacaagct gtgcgaccag gtgtcggacg ctgtgctgga 180
 ctctgcctg ggcgaggacc ccgacagcaa agtggcctgc gagacctgca ccaagacgaa 240
 catggtgatg gtgttcggcg agatcancan caangcgagc gtggactacg agaagatngt 300
 gcgcgacacc t 311

<210> 225
 <211> 293
 <212> nucleic acid
 <213> Zea mays

 <400> 225

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 aaggagaact togacttcag gcccgggatg atcagcatca acctcgacct gaagaagggc 120
 ggcaacaggt tcatcaagac cgccgcctac ggccacttcg gccgtgacga cgccgacttc 180
 acctgggagg tgggtgaagcc cctcaagttc gacaaggcat cggcttaagg ttgggagtgt 240
 cactgtggac atgaggacta ccttcctctg gctctgctgt tacctgcaag cat 293

<210> 226
 <211> 225
 <212> nucleic acid
 <213> Zea mays

 <400> 226

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 gatcctcaag atcgtgaagg agaacttcga cttcaggccc gggatgatca gcatcaacct 120
 cgacctgaag aagggcgcca acaggttcat caagaccgcc gcctacggcc acttcggccg 180
 tgacgacgcc gacttcacct gggaggtggt gaagcccctc aagtt 225

<210> 227
 <211> 256
 <212> nucleic acid
 <213> Zea mays

 <400> 227

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 ggccggaaga tcatcatcga cacctacggt ggctggggag cccacggcgg gggcgccctc 120
 tccggcaagg acccgaccaa ggtggaccgc agcggggcct acgtcgcgag gcaggctgcc 180
 aagagcatcg tcgccgccgg cctcgcccgc ngtgccatcg tcaagtnttc naaggcatcg 240
 ggttgcccga anccct 256

<210> 228

<211> 281
 <212> nucleic acid
 <213> Zea mays

 <400> 228

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 nacccaacca aggttgaccg cagcggagcc tatgtcgcaa ggcaggctgc caagagcatc 120
 gtcgccagcg gccttgctcg ccgcgccatc gtccaggtgt cttacgccat cggntgcccg 180
 agcctctctc cgtgttcgtc gacacgtacg gcaccggcgc gatccccgac aaggagatct 240
 caagattgtc aaggagaatt cgatttcagg ctggcatgat c 281

<210> 229
 <211> 290
 <212> nucleic acid
 <213> Zea mays

 <400> 229

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 atccagagaa gatggcagct gtcgacacat tctctttcac ctccgagtct gtgaacgagg 120
 gacaccctga caagctctgt gaccaggtct cagatccgtt cttgacgctt gccttgctga 180
 ggaccctgac agcaagggtg cttgtgagac ctgcaccaag accaacaatgg tcatggtctt 240
 tggtgagatc accaccaagg ccaatgttga ctacgagaag attgtcaggg 290

<210> 230
 <211> 318
 <212> nucleic acid
 <213> Zea mays

 <400> 230

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 gaagaaggca atggcggccg agagcttcc tttcacctcg gagtccgtga aacgaggggc 120
 accccgaaca agctgtgcga ccangtgtcg gacgccgtgc tttgacgcat gnctcncgca 180
 ngagnccgac agcaangtgg cctgcgagac ctgcaccaag ancaacaatgg tgatggtggt 240
 cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacctgccg 300
 cgagatcggg ttcactcc 318

<210> 231
 <211> 276
 <212> nucleic acid
 <213> Zea mays

 <400> 231

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 accaaggttg accgcagcgg agcctatgtc gcaaggcagg ctgccaagag catcgtcgcc 120
 agcggccttg ctgcgcgcgc catcgtccag gtgtcttacg ccatcggttg cccgagcctc 180
 tctccgtgtt cgtcgacacg tacggcaccg gcgcgatccc cgacaaggag atcctcaaga 240
 tgtcaaggag aattcgattt cangctggca tgatca 276

<210> 232
 <211> 244
 <212> nucleic acid
 <213> Zea mays

 <400> 232

 ctcaaggcga tgctggcctc actggccgca agatcatcat tgacacctac ggtggctggg 60
 gagcccatgg tgggtggcgct ttctccggca aggacccaac caaggttgac cgcagcggag 120
 cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct cgccgcgcca 180
 tcgtccaggt gtcttacgcc atcgggtgcc cgagcctctc tccgtgttcg tcgacacgta 240
 cggc 244

<210> 233
 <211> 349
 <212> nucleic acid
 <213> Zea mays

 <400> 233

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 tgagcagcag tcccctgata ttgctcaggg tgtgcatggc cacttcacca agcgcgccga 180
 ggagattgga gctggtgacc agggacacat gttcgggtat gcgaccgatg agaccctgag 240
 ttgatgcctt caagccatgt cttgccacca gctagtgtc gtctcacgag gccgcaaaan 300

ggactgnccn ggnnaanctg atggagacca gtganatcga gtacncatt 349

<210> 234
 <211> 228
 <212> nucleic acid
 <213> Zea mays

<400> 234

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ccatggtggt ggcgctttct ccggcaagga cccaaccaag gttgaccgca gcggagccta 120

tgtcgcaagg caggctgccca agagcatcgt cgccagcggc cttgctcgcc gcgccatcgt 180

ccagggtgtct tacgccatcg gcgtgcccga gcctctctcc gtgttcgt 228

<210> 235
 <211> 263
 <212> nucleic acid
 <213> Zea mays

<400> 235

cctgacagca aggttgcttg cgagacctgc accaagacca acatggtcat ggtcttttgt 60

gagatcacca ccaaggccaa tgtcgactac gagaagattg tcagggagac atgccgcaac 120

atnpgtttctg tgtcgaaaga tgtcgggctt gacgctgacc actgcaagggt gcttgtnaac 180

attnagcagc agtccccctga tattgctcag ggtgtgcagg ccattcacna agccccccga 240

ggagntngag tgtgaccagg gga 263

<210> 236
 <211> 219
 <212> nucleic acid
 <213> Zea mays

<400> 236

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ggcanccccga caagctgtgc gaccaggtgt cggacgccgt gcttgacgca tgcctcgcgc 120

aggacccccga cagcaagggtg gcctgcgaga cctgcaccaa gaccaacatg gtgatgggtgt 180

tcggcgagat cacgaccaag gcgaccgtgg actacgaga 219

<210> 237

<211> 301
 <212> nucleic acid
 <213> Zea mays

 <400> 237

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 tggtagagcg agcgagaaga aggcaatggc ggcgagagc ttctgttca cctcggagtc 120
 cgtgaacgag gggcaccag acaagctgtg cgaccaggtg tcggacgcgg tgctggacgc 180
 ctgcctggcg caggaccccg acagcaaggt ggctgcgag acctgcacca agacgaacat 240
 ggtgatggtg ttggcgaga tnaacaaca aggcgagcgt ggactacgag aagatggtgc 300
 g 301

<210> 238
 <211> 439
 <212> nucleic acid
 <213> Zea mays

 <400> 238

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 gcgcgcgcgc agatcaaaga agatggcagc tgctgcacaca ttctcttca cctcggagtc 180
 tgtgaacgag ggacaccctg acaagctctg tgaccaggtc tcagatgccg ttcttgacgc 240
 ttgccttgct gaggaccctg acagcaaggt tgcttgtgag acctgcacca agaccaacat 300
 gggcatggnc tttggtgaga tcaccaccaa ggccaatggt gactaccaag aagattgtca 360
 gggagnacct gccgnaacat tnggtttggg gtcaaacgat gttggcttga cgccaaccac 420
 tggaaggggc tcgtnaant 439

<210> 239
 <211> 239
 <212> nucleic acid
 <213> Zea mays

 <400> 239

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 tctctatctc caccagcac gacgagacag tgaccaatga tgagatcgct gctgacctga 120

aggagcatgt catcaagcct gtcacccctg agcagtagct tgacgagaag accatcttcc 180
accttaaccc atctggccgc nttgtcattg gtggactcac ggcgatgtgg cctcactgg 239

<210> 240
<211> 224
<212> nucleic acid
<213> Zea mays

<400> 240

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cactgcaagg tgcttgtgaa cattgagcag cagtccctg atattgtca ggggtgtgcac 120
ggccacttca ccaagcgccc cgaggagatt ggagctgggt accaggggca catgtttngg 180
tatnctantt acnngacacc tgagctgatg ccctcagcc atgt 224

<210> 241
<211> 274
<212> nucleic acid
<213> Zea mays

<400> 241

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aagatggccg gactcgacac ctctctcttc acctcggagt ccgtgaacga gggacaccct 120
gacaagctct gcgaccaggt ctcatatgtt gttctggacg cttgccttgc tgaggaccct 180
gacagcaagg ttgcttgcca gacctgcacc aagaccaaca tggatcatgt ctttggtgag 240
atcaccacca aggccaatgt cgatacgaga agat 274

<210> 242
<211> 232
<212> nucleic acid
<213> Zea mays

<400> 242

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tgagatcacc accaaggcca angtcgacta cgaggagatt gtcagggaga catgccgcaa 180
cattggtntc gtgtcgaacg atgtcgggct tgacgctgac cactgcaagg tg 232

<210> 243
 <211> 267
 <212> nucleic acid
 <213> Zea mays

 <400> 243

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 agaaggcaat ggcgggccgag agcntccttt tcacctcgga gtccgtgaac gaggggcacc 120
 ccgacaagct gtgcgaccag gtgtcggacg ccgtgcttga cgcctgcctc gcgcaggacc 180
 ccgacagcaa ggtggcctgc gagacctgca ccaagaccaa catggtgatg gtgttcggcg 240
 agatcacgac caaggcgacc gtggact 267

<210> 244
 <211> 309
 <212> nucleic acid
 <213> Zea mays

 <400> 244

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 ccagcttgcc ccaggttggt agagcgagcg agaagaaggc aatggcgggcg gagagcttcc 120
 tgttcacatc ggagtccgtg aacgaggggc acccagacaa gctgtgcgac caggtgtcgg 180
 acgcggtgct ggacgcctgc ntggcgagg accccgacag caaggtggcc tgcgagaccn 240
 gcaccaagac gaacatggtg atggtgttcg gcgagatcac caccaaggcg agcgtggaca 300
 acgagaaga 309

<210> 245
 <211> 347
 <212> nucleic acid
 <213> Zea mays

 <400> 245

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 caaagaagat ggcagctgtc gacacattcc tcttcacctc ggagtctgtg aacgagggac 120
 accctgacaa gctctgtgac caggtctcag atgccgttct tgacgcttgc cttgctgagg 180
 accctgacag caaggttgct tgtgagacct gnaccaagac caacatggtc atggtctttg 240

gtgagatcac caccaaggcc attgtgacta cgagaagatt gtccaggag acttgccgca 300
acattgggttt gtgtcaaacg atgttgggnt gacgccacac tgcaaag 347

<210> 246
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 246

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tganctgac cactgcaagg tgcttgtaaa cattggnnca gcagtcccct natattgctc 120
agngtgtgca cggccacttc accaagcgcc ccgaggagat tggagctggt gaccaggggc 180
acatgttttg gtatgcnact gacgagaccc ctgagctgat gcccctcagc catgtccttg 240
ccaccaagct tgggtgctcgt c 261

<210> 247
<211> 211
<212> nucleic acid
<213> Zea mays

<400> 247

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tggcctgcga gacctgcacc aagaccaaca tggatgatgt gttcggcgag atcacgacca 120
aggcgaccgt ggactacgag aagatcgtgc gcgacacctg ccgagagatc gggttcacct 180
ccgacgacgt gggctcgacg ccgaccgctg c 211

<210> 248
<211> 301
<212> nucleic acid
<213> Zea mays

<400> 248

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cccgacaagg aganncggn ganngtcaag gagaacttcg acttcaggcc agggatgatc 120
accatcaacc tcgacctcaa gaaggcggc aacaggttca tcaagaccgc cgcatacggc 180
cactttggcc gtgacgacgc cgacttcacc tgggagngg tcaagcccct aaagaaggca 240

tccgcttaag aatgtattgg gaagttcact ggacatgagg ttcattctcg tctggctctg 300

c 301

<210> 249

<211> 320

<212> nucleic acid

<213> Zea mays

<400> 249

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gcccggcatg atcatcatca acctcgacct caagaaaggc ggcaacgggc gctacctcaa 120

gacggcgggc tacgggcact ttgggaggga cgaccccgac ttcacctggg aggtgggtgaa 180

gcccctcaag gcggagaagc cgtcttctgc atgaggcgcc tcctctgttt tggaagaagc 240

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ctacttgtga ttcttgatct 320

<210> 250

<211> 282

<212> nucleic acid

<213> Zea mays

<400> 250

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gccaccgacg agacccccga gctgatgccg ctgagccacg tgctggccan caagctgggc 120

gcgcgctcac cgagncacgc aagaacggca ctgcgcgtgg ctgagccacn gacggcaaga 180

cccaggtgac ggtggagtac gtgaacgatg gcggcgccat ggtgcccgtc cgcggtgcaca 240

cgtgctcacc tncaccacgc acgacganna ctnaacaacg ac 282

<210> 251

<211> 239

<212> nucleic acid

<213> Zea mays

<400> 251

cttcacctc aaccgctctg gccgcttcgt catcgcgga cctcacggcg acgctggcct 60

cactggccgg aagatcatan tccaggtctc ctacgccatc ggcgtgcccg aaaccctgt 120

cggtgttcgt ggacacgtac ggcaccggcg cgatccccga caaggagatc ctgaagatcg 180
tgaaggagaa cttcgacttc aggcccggca tgatcatcat caacctcgac ctcaagaaa 239

<210> 252
<211> 511
<212> nucleic acid
<213> Zea mays

<400> 252

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cgtccaccac ggnggtggcg ctttctccgg caaggacccc accaaggtgg accgcagcgc 120
gcctacgtgg ccaggcaggc cgccaagagc atcgtggcca gcggcctcgc ccgcgcgcgc 180
tcgtgcaggt gtcgtacgcc atcggcgtgc cggagcccct gtccgtgttc gtcaaccgta 240
cggcaacggc acgattcccc acaagggaga ttctcaaaga tcgttaagga gaaactcgaa 300
cttnaanggc cgggaanatt aagcaatcaa acctncgaac ctngaaaaaa gggcggcaac 360
aaggttnaat caaaaancgc ccgcctaagg ncaattnggc ccgttaacca acgcgantt 420
aactgggaag tggtnaaacc cctcaaattc cacaaagcat cggcttaaag gttggaattt 480
cactgtggac attaaggact aancttctc t 511

<210> 253
<211> 234
<212> nucleic acid
<213> Zea mays

<400> 253

cnacngtctn gaatgagcac gtcancaagc ccgtcatccc ggagaggtac ctggacgaga 60
agacnatctn ccnnnnnnnn ncgtcnnggc gtttcgtcat cggcgggccc cacggggacg 120
ccggcctcac cggccgcaag atcatcatcg acacctacgg cggctgggga gccacggcg 180
ggggcgccct ctccggcaag gacccaccca angtggaccg cngcggggcc tact 234

<210> 254
<211> 295
<212> nucleic acid
<213> Zea mays

<400> 254

cttttctcct cctcctgnc gggtccttaa taaagagcag cagcgcaagg tgagccgcca 60
 gcttgcccca ggttggtaga gcgagcgaga agaaggcaat ggcggcggag agcttcctgt 120
 tcacgtcgga gtccgtgaac gaggggcacc cagacaagct gtgcgaccag gtgtcggacg 180
 cgggtgctgga cgctgcctg ggcgaggacc ccgacagcaa ggtggcctgc gagacctgca 240
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<210> 255
 <211> 257
 <212> nucleic acid
 <213> Zea mays
 <400> 255

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 gaccncacgg cgatgcnggc ctacnnggc gcaagancan catngacacc nacggtggcn 180
 ggggagccca tgggtggtggc gcntncnccg gcaaggaccc aaccaagggt gaccgcagcg 240
 gacctatgtc gcaaggc 257

<210> 256
 <211> 206
 <212> nucleic acid
 <213> Zea mays
 <400> 256

ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga cacctgccgc 60
 gagatcgggt tcacctccga cgacgtgggc ctgcagccg accgtgcaa ggtgctggtg 120
 aacatcgagc agcagtcacc cgacatcgcg cagggcgctgc acgggcactt cacgaagcgg 180
 cccgaggaga tcggcgcggg cgacca 206

<210> 257
 <211> 208
 <212> nucleic acid
 <213> Zea mays
 <400> 257

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ctggcctcac tggccggaag atcatcatcg acacctacgg tggctgggga gcccacggcg 120
 ggggcgcctt ctccggcaag gacccgacca aggtggaccg cagcggggcc tacgtcgcga 180
 ggcaggctgc caagagcatc gtcgccgc 208

<210> 258
 <211> 339
 <212> nucleic acid
 <213> Zea mays
 <400> 258

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 gtcatccccg agcagtacct cgacgagaag acaatcttcc acctcaaccg gtctggccgt 180
 tcgtcatcgg cggaacctac ggcgacgcgg cctcactggc ggaagatcat catcgacacc 240
 tacgggtgntt gggagccacg gcggggcgct ttncggcaag gaccgncaag tggacgancg 300
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<210> 259
 <211> 195
 <212> nucleic acid
 <213> Zea mays
 <400> 259

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 ttcggcgaga tcacgaccaa ggcgaccgtg gactacgaga agatcgtgcg cgacacctgc 120
 cgcgagatcg ggttcacctc cgacgacgtg ggctcgacg ccgaccgctg caaggtgctg 180
 gtgaacatcg agcag 195

<210> 260
 <211> 267
 <212> nucleic acid
 <213> Zea mays
 <400> 260

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 agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcgag tccgtgaacg 120

aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
 cgcaggaccc cgacagcaag gtggcctgcg agacctgcac caagaccaac atgggtgatgg 240
 tgttcggcga natcacgacc aaggcga 267

<210> 261
 <211> 272
 <212> nucleic acid
 <213> Zea mays

<400> 261

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 acgaggggca ccccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc 180
 tcgogcagga ccccgacagc aagggtggcct gcgagacctg caccaagacc aacatgggtga 240
 tngtgttcgg cgagatcacg accaangcga cc 272

<210> 262
 <211> 335
 <212> nucleic acid
 <213> Zea mays

<400> 262

ggggactaat gtgcgaaggc aggctgccaa gagcatcgtc gccagcggcc ttgctcgccg 60
 cgccatcgtc cagggtgtctt acgccatcgg gtgcccagagc ctctctccgt gttcgctgac 120
 acgtacggca ccggcgcgat ccccgacaag gagatcctca agattgtcaa ggagaactcg 180
 atttcaggcc tggcatgac atcatcaacc ttgacctcaa gaaaggcggc aacggggcgct 240
 actcaagagn gcggctacgg ccactttgga agggacgacc tgattcacct gggaggtggt 300
 gaagccattc aatcgagaa actttgctaa gcggc 335

<210> 263
 <211> 270
 <212> nucleic acid
 <213> Zea mays

<400> 263

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 cgtgaacgag gggcaccgac acaagctgtg cgaccagggtg tcggacgccc tgcttgacgc 180
 atgcntcgcg caggaccgac acagcaaggt ggctgagagc acctgcacca agaccaacat 240
 ggtgatggtg ttccggcgaga tcacgaccaa 270

<210> 264
 <211> 246
 <212> nucleic acid
 <213> Zea mays
 <400> 264

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 tctgtgacca ggtctcagat gccgttcttg acgcttgccct tgctgaggac cctgacagca 180
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 ccaagg 246

<210> 265
 <211> 263
 <212> nucleic acid
 <213> Zea mays
 <400> 265

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 cnccgacaag ctgtgagacc aggtgtcgga cgccgtgctt gacgcattgcc tcgagcagga 180
 ccccgacagc aaggtggcct gcgagacctg caccaagacc aacatggtga tgggtgttcgg 240
 cgagatcacg accaaggcga ccg 263

<210> 266
 <211> 295
 <212> nucleic acid
 <213> Zea mays
 <400> 266

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aaagaagatg gcagctgtcg acacattcnc tcttcacctn nggagtctgt gaacgagggg 120
 caccctgaca agctctgtga ccaggtctca gatgccgttc ttgacgcttg ccttgctgag 180
 gaccctgaca gcaaggttgc ttgtgagacc tgcaccaaga ccaacatggt catggtcttt 240
 ggtgagatca ccaccaaggg caatgttgac tacgagaaga ttgtcagggg gacct 295

<210> 267
 <211> 288
 <212> nucleic acid
 <213> Zea mays
 <400> 267

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 ttgtgaacat tgagcagcag tcccctgata ttgtcaggg tgtgcacggc cacttcacca 120
 agogccccga ggagattgga gtggtgacca ggggcacatg tttgggtatg cgactgacga 180
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 agttcgccaa gaatggaact gccctggcna agcccgatgg aagaccag 288

<210> 268
 <211> 254
 <212> nucleic acid
 <213> Zea mays
 <400> 268

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 tggaccgcag cggggcctac gtcgcgaggg aggctgcaa nagcatcgtc gccgcccggc 180
 taaccgcgcg cgnctttgtc caggtctnct acgccatcgg ntnaccgag cccctttcgg 240
 tgttcgtgga cagc 254

<210> 269
 <211> 525
 <212> nucleic acid
 <213> Zea mays
 <400> 269

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tcgacccacg cgtccgccc cgcgtccgcc cagcgcgcgc cccacgcgctc cgcccacgcg 120
tccgggtcgga tctgggacga gacgagacga tcccngcctc cctcaaccg gaacttgttt 180
taccatct catccactg actccacca cccaccgcc cgtgcctcc gccggatctc 240
gtcggactcg gatccgccc accacgacca cccgcgttg ncgcgcgca gcagcagcag 300
atcagagaag atggccggac tcgacacctt cctcttcacc tcggagtcgc tgaacgaggg 360
acaccctgac aagctctgcg accaggtctc agatgctgtt ctggacgctt gccttgctga 420
ggaccctgac agcanggttg cttgcgagac ctgcaccaag accaaccatgg tcatggcttt 480
ggtgagatca ccaccaagn caatgntgac tttttnaana ntttg 525

<210> 270
<211> 312
<212> nucleic acid
<213> Zea mays

<400> 270
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ctcaagaaag ggcgcaaccg gcgctacctc aagacggcgg cctacgggca ctttgggagg 120
gacgaccccg acttcacctg ggaggtggtg aagcccctca aggcggagaa gcgctcttct 180
gcatgaggcg cctcctctgt tttggaagaa gcttttggtc tggctctggtc tggctctggtg 240
tgctgcgct ctatcatgct ttttatggc tctacttgt gattcttgat ctgccccttg 300
cttatcattg ta 312

<210> 271
<211> 227
<212> nucleic acid
<213> Zea mays

<400> 271
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ggcnaentc gcgaggcagn ctgccaanag catcgtcgcc gccggcctcn cncgcgcgc 120
cattgtccag gtctcctacg ccatcggcgt gccgagccc ctttcggtgt tcgtggacac 180
gtacggcacc ggcgcgatcc ccgacaagga gatcctgaag ancgtagg 227

<210> 272

<211> 234
 <212> nucleic acid
 <213> Zea mays

 <400> 272

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 ttcacctcgg ngtcctgtaa cgagggggcac ccagacaagc tgtgcgacca ggtgtcggnc 120
 ggggtgctgg acgcctgcct ggcgaggac cccgacagca aggtggcctg cgagacctgc 180
 accaagacga acntngtgat ggtgttcggc gagatcncca ccaaggcgag cgtg 234

<210> 273
 <211> 239
 <212> nucleic acid
 <213> Zea mays

 <400> 273

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 caacatcggt tgacacaaaa ccaatgttgc ggcaggtctc cctgacaatc ttctcgtagt 120
 caacattggc cttggtggtg atctcaccaa agaccatgac catgttggtc ttggtgcagg 180
 tctcacaagc aaccttgctg tcagggtcct cagcaaggca agcgtcaaga acggcatct 239

<210> 274
 <211> 245
 <212> nucleic acid
 <213> Zea mays

 <400> 274

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 gcaatggcgg ccgagagctt ctttttcacc tcggagtccg tgaacgaggg gcaccccgac 120
 aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180
 agcaaggtgg cctgcgagac ctgcaccaag accaacaagg tgatggtggt cggcgagatc 240
 acgac 245

<210> 275
 <211> 268
 <212> nucleic acid
 <213> Zea mays

<400> 275

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agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
cgcaggaccc cgacagcaag gtggcctgcg agacctgcac caagaccaac catggtgatg 240
gtgttcggcg agatcacgac caaggcga 268

<210> 276

<211> 228

<212> nucleic acid

<213> Zea mays

<400> 276

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gagcccatgg tggcggcgct ttctccggca aggaccaac caaggctgac cgcagcggag 120
cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct cgcgcgcga 180
tcgtccaggt gtcttacgcc atcgggtgcc cgagcctctc tccgtggt 228

<210> 277

<211> 253

<212> nucleic acid

<213> Zea mays

<400> 277

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cgagaagacc atcttcacc ttaaccatc tggccgcttt gtcattggtg gacctcacgg 120
cgatgctggc ctactggcc gcaagatcat cattgacacc nacggtggct ggggatcccn 180
nnggggtggc cttttctcgg aagagggcna aacnaagggt gncgtagtgg tntttntga 240
aanggtagnn tgc 253

<210> 278

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 278

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 ttggttagagc gagcgagaag aaggcaatgg cggcggagag cntcctgttc acctcggagt 120
 ccgtgaacga ggggcaccca gacaagctgt gcgaccaggt gtcggacgcg gtgctggacg 180
 cctgcctggc gcaggacccc gacagcaagg tggcctgcga gacctgcacc aagacgaaca 240
 tggatgatggg gttcggcgag atcaccac 268

<210> 279
 <211> 218
 <212> nucleic acid
 <213> Zea mays

<400> 279

cccgaccaag gtggaccgca gcggggccta cgtcgcgagg caggctgcca agagcatcgt 60
 cgccgcgggc ctgcgccgcc gcgccatcgt ccaggctctcc tacgccatcg ggtgcccgag 120
 cccctatcgg tgttcgtgga cacgtacggc ancggcgcga tccccgacaa ggagatcctg 180
 aagatcgtga aggagaactt cgacttcaag cccggcat 218

<210> 280
 <211> 314
 <212> nucleic acid
 <213> Zea mays

<400> 280

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 cggccgagag ctctcttttc accttcggag tccgtgaacg aggggcaccc cgacaagctg 120
 tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc cgacagcaag 180
 gtggctgcga gactgcacaa gaccaacatg gtgatgggtg tcggcgagat cacgaacaan 240
 gcgacgtgga ctacgagaag atcgtgcgcg acacctgccc gcgagatcgg gttcacctcc 300
 gacgacgtgg gctc 314

<210> 281
 <211> 216
 <212> nucleic acid
 <213> Zea mays

<400> 281

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 caagagcatc gtcgccgccc gctcgccc cgcgccatt gtccaggtct cctacgcca 120
 cgggtgcccg ancccccttc ggtgttcgtg gacacgtacg gcaccggcgc gatccccgac 180
 aaggagatcc tgaagatcgt gaaggagaac ttcgac 216

<210> 282
 <211> 289
 <212> nucleic acid
 <213> Zea mays
 <400> 282

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 attgacacct acggtggctg gggagcccat ggtggtggcg cttctccgg caaggaccca 120
 accaaggttg accgcagcgg acctatgtcg caaggcaggc tgccaagagc atcgtcgcca 180
 ggggecttgc tcgccgcgcc atcgccagn tgtcttacgc canngggtgc nngancctct 240
 ctccgtgttc gaaaanannn anngcnngn nntcccccaa nggttttct 289

<210> 283
 <211> 247
 <212> nucleic acid
 <213> Zea mays
 <400> 283

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 acaagctgtg cgaccaggtg tcggacgccg tgcttnangc atgcctcgcg caggacccccg 180
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 tcacgac 247

<210> 284
 <211> 275
 <212> nucleic acid
 <213> Zea mays
 <400> 284

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agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
 aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
 cgcaggaccc cgacagcaaa gttgctgca aactgcacc aagancaaca tggatgatgt 240
 gttcggcgag atcacgacca aggggaccgt ggatt 275

<210> 285
 <211> 255
 <212> nucleic acid
 <213> Zea mays
 <400> 285

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 cgagggggcac ccagacaagc tgtgcgacca ggtgtcggac gcggtgctgg acgcctgcct 180
 ggcgaggagc cccgacagca aggtggcctg cgagacctgc accaagacga acatggtgat 240
 ggtgttcggc gagat 255

<210> 286
 <211> 221
 <212> nucleic acid
 <213> Zea mays
 <400> 286

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 ggctacgtc ncnaggcagg cngccaagag catcgtcncc nccggcctcg ccnccgcnc 120
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 tacggcaccg gcgcgatccc cnacaaggag atcctgaaga t 221

<210> 287
 <211> 216
 <212> nucleic acid
 <213> Zea mays
 <400> 287

ctctaccag cacgacgaga cagtcaccaa cgacgagatt gctgctgacc tgaaggagca 60
 cgtcacaaag ccagtcaccc ccgagcagta cctcgacgag aagacaatct tccacctcaa 120

cccgctctggc cgcttcgtca tcngcggacc tcacggcgac gctggcnctnn ctgnnnccggn 180
agatcatcat cgacacctan ggttgctggg gagcca 216

<210> 288
<211> 292
<212> nucleic acid
<213> Zea mays

<400> 288

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tcaaggagaa cttcgatttc aagcctggca tgatcatcat caacottgac ctcaagaaag 120
gcggaacgg gcgctacctc aagacggcgg cctacggcca ctttgaagg gacgacctg 180
acttcacctg ggaggtggtg aagccacttc aagtcggaga aacottctgc ctaaggcggc 240
ctttttttca gtaagaagct tttggtggtc tgctgtgctt aatcagcttt ta 292

<210> 289
<211> 276
<212> nucleic acid
<213> Zea mays

<400> 289

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aacgggcgct acctcaagac ggcggcctac gggcactttg ggagggacga ccccgacttc 120
acctgggagg tggatgaagcc cctcaaggcg gagaagccgt cttctgcatg aggcgcctcc 180
tgtgtttcgg aagaagcttt tggctctggtc tgccctgcgt ctatcatgct tttttatggc 240
tctacgtgt tgtgattctt gatctgcccc ttgctt 276

<210> 290
<211> 219
<212> nucleic acid
<213> Zea mays

<400> 290

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ccccgacaag gagatcctca agatcgtgaa ggagaacttc gacttcangc ccgggatgat 120
cagcatcaac ctcgacctga agaaggcgg caacaggttc atcaagaccg ccgcctacgg 180

ccattcggcc gtgacgacgc cgacttcact gggagtgggt 219

<210> 291
 <211> 191
 <212> nucleic acid
 <213> Zea mays

<400> 291

catgccgcaa cattggtttc gtgtcgaacg atgtcgggct tgacgctgac cactgcaagg 60
 tgcttgtgaa cattgagcag cagtcccttg atattgctca ggggtgtgcac ggccacttca 120
 ccaagcgccc cgaggagatt ggagctgggtg accaggggca catgtttggg tatgcgactg 180
 acgagacccc t 191

<210> 292
 <211> 315
 <212> nucleic acid
 <213> Zea mays

<400> 292

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 actccgcctc gaccggnnncn cgtcggactc gganccgccc gaccaccccg cgccgcccga 120
 gatcaaagaa gatggcagcn gtcgacacat tctctttcac ctcgaggtct gtgaacgagg 180
 gacaccctga caagcncctgt gaccaggtct cagatgccgt tcttgacgct tgccttgngg 240
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 ttggtgagan nacca 315

<210> 293
 <211> 501
 <212> nucleic acid
 <213> Zea mays

<400> 293

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 caaggagatc ctcaagatcg tcaaggagaa cttcgacttc aggccccggg atnatcacca 180
 tcaacctcga cctcaanaaa gggcgggaac aggttcatca agaccgccgc atacngncac 240

tttggcccgt gacgancna ctttacctgg gaagtgggtca atccccctaaa gaaagccatn 300
ccncttaaga atgtanttgg naagtttact tggacatgaa gttcattctt ngctctngctt 360
ctgctgatnc cctnnaanga ttgcttgntn cttgcttgcc cctngattgt ntgttttgan 420
caantgantt ngcttgntct tgttccatnt gaaaaaccnn attaatngtg gncnttttgg 480
tgaaaaaaaag ntttngccna t 501

<210> 294
<211> 281
<212> nucleic acid
<213> Zea mays
<400> 294

gcctctttctc ctccctcctg ccgggtcctt aataaagagc agcagcgcaa gaggttggtta 60
gagcgagcga gaagaaggca atggcgggcg agagcttcct gttcacctcg gagtccgtga 120
acgaggggca cccagacaag ctgtgcgacc aggtgtcgga cgcggtgctg gacgcctgcc 180
tggcgagaa ccccgacagc aagtggcctg cgagacctgc accaagacga acatggtgat 240
ggtgttcggc gaantcacca ccagggnagn tggatacgaa t 281

<210> 295
<211> 486
<212> nucleic acid
<213> Zea mays
<400> 295

gggnanttnt tnnagncctt cnacgcgnnc agtaccggtc acagaattcc cgggncgacc 60
acgcgtccnc ggacgcgtgg gcgacaagga gatcctcang atcgtgaagg agannttcac 120
ttcaggcccg ggatgatcag catcaacctc gacctgaaga agggcgga caggttctca 180
agaccgccgc ctacggccac ttcgggcgtg acnacgccga cttcacctgg gaggtgtgaa 240
gcccctcaag ttogacaagg catcggctta aggttgggan tgtcactgtg gacataggac 300
taccttctc tggctctgct gttacctgca agcattgctg ctgctggatg tgtggtttga 360
tcagtgaactg gctgctgctc catagaagat gaacggagag aaggatgatg aangctttgg 420
caatcgcccg ctgcaactgc aacctatgcc atgccggctt aatgattggg taaattttgg 480
cttnca 486

<210> 296
 <211> 173
 <212> nucleic acid
 <213> Zea mays

<400> 296

tacgtcgcca ggcaggccgc caagagcatc gtggccagcg gcctcgcccg ccgctgcctc 60
 gtgcaggtgt cctacgccat cggcgtgccg gagcccctgt ccgtgttcgt cgactcctac 120
 ggcaccggga ccatccccga caaggagatc ctaaagatcg tcaaggagaa ctt 173

<210> 297
 <211> 278
 <212> nucleic acid
 <213> Zea mays

<400> 297

cgctctttct cctccctcct gccgggtcct taataaagag cagcagcgca agaggttggt 60
 agagcgagcg agaagaaggc aatggcgccg gagagcttcc tggtcacctc ggagtccgtg 120
 aacgaggggc acccagacaa gctgtgcgac cagggtgcgg acgcggtgct ggacgcctgc 180
 ntggcgagg accccgacag caaggtggcc tgcnatncct ncacnangac gaacatgggtg 240
 atggtgttcg gcgaaatcac cncnantgcg acntngac 278

<210> 298
 <211> 309
 <212> nucleic acid
 <213> Zea mays

<400> 298

ggcgtgccgg agcccctgtc cgtgttcgtc gactcctacg gcaccgggac catccccgac 60
 aaggagatcc taaagatcgt caaggagaac ttcgacttca ggccagggat gatcaccatc 120
 aacctcgacc tcaagaaggc cggaacagg ttcatacaaga ccgccgcata cggccacttt 180
 ggcntgacga cgccgacttc acctgggagg tggtaagcc cctaaagaag gcatccgctt 240
 aagaatgtat tgggaagtgc actggacatg aggttcatct tcgtctggct ctgctgatac 300
 ctgcaagat 309

<210> 299

<211> 197
 <212> nucleic acid
 <213> Zea mays

 <400> 299

 gggagaccng ccgcaacatt ggttttgtgt caaacgatgt tgggcttgac gccgaccacn 60
 gcaaggtgct cgtgaacatt gagcagcagt cccctgatat tgctcagggt gtgcatggcc 120
 acttcaccaa gcgccccgag gagattggag ctggtgacca gggacacatg ttcgggtatg 180
 cgaccgatga gaccctt 197

<210> 300
 <211> 253
 <212> nucleic acid
 <213> Zea mays

 <400> 300

 cgtttgcctc ttctccctct tgccggctcc gaataaagag cagcagcgca agaggctcgg 60
 agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
 aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180
 cgcaggaccc cgacagcaag gtggctgcga gacctgcacc nagaccaaca tggatgatngt 240
 gttcggcgag atc 253

<210> 301
 <211> 302
 <212> nucleic acid
 <213> Zea mays

 <400> 301

 cgtgaaacga ggggacaccc tgacaagctc tgcgaccagg tctcagatgc tgttctggac 60
 gcttgccctg ctgaggaccc tgacagcaag ttgcttgag acctgcacca agaccaacat 120
 ggtcatggtc tttggtgaga tcaccaccaa ggccaatgtc gactacgaga agattgtcag 180
 gggagacatg ccgcaaaatt ggtttcngt naganngatg tggggcttga ngntgaccac 240
 ngcanggtgn tnatanaaan ttngnagnan tncctgtat ttgcgcnngg gggnnncncc 300
 nc 302

<210> 302

<211> 460
 <212> nucleic acid
 <213> Zea mays

 <400> 302

 cgccgcccgc ctgcgccgcc gcgccatcgt ccaggtctcc tacgccatcg gcgtgcccga 60
 gcccctatcg gtgttcgtgg acacgtacng gancngngcn anncccgacn agggnaatcn 120
 tganaatngn anaagganaa nttnanantn caggcccggt tgatcattat naacctagac 180
 ctcaanaaag gcggaaacgg gcnctaccta aagacggggg tctacgggcn ctttgngagg 240
 gacgaccgag anttcacctg agaggtggna aagcccctca aggcggaaaa gccgtcttct 300
 gcatgaggcg cctcctctgt ttengaagaa gcttttggtc tggctctgct gcgctctatc 360
 atgctttttt atggctncta cgtgttggtga ttcttgatct gcccttgct tatcatttgt 420
 actgtactgt cactgtccta ataagtggta cgtgtgcggg 460

<210> 303
 <211> 297
 <212> nucleic acid
 <213> Zea mays

 <400> 303

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 natngnceca ggtgtattat cgccatccgg cnngncngng cntctcnccg tgttcgtcga 120
 cacgtacggc accggcgcga tccccgacaa gngnncnct gaagattgta caaggagaac 180
 ttcgatttca ggctggcat gatcatnnnc aaccttganc tcaagaaagg nggcaacggg 240
 cgctacctca agacggcggc ctacggcnac tttggaaggg acgaccctga cttcacc 297

<210> 304
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 304

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 ctcatcncaa ctccgtctca ccgaggtccg caagaacgga acctgccct ggctcaggcc 120
 tgatgggaag acccaggtga cagtcgagta ccgcaatgag ggtggtgcc tggccccat 180

ccgtgtccac accgtctca tctccacca gcacgacgag acagtgacca atgatgagat 240
cgctgctgac ctgaaggag 259

<210> 305
<211> 244
<212> nucleic acid
<213> Zea mays

<400> 305

cttctctctcc ctctctgccg gtccttaata aagagcagca gcgcaagagg ttggtagagc 60
gagcgagaag aaggcaatgg cggcggagag cttcctgttc acctcggagt ccgtgaacga 120
ggggcacca gacaagctgt gcgaccaggt gtcggacgcg gtgctggacg cctgcctggc 180
gcaggacccc gacagcaagg tggcctgcga nacctgcacc aagacgaaca tggatgatgt 240
gttc 244

<210> 306
<211> 236
<212> nucleic acid
<213> Zea mays

<400> 306

cgctggactc ggatccgccc gaccacgacc accccgcgcc gccgccgccc acagcagcag 60
atcagagaag atggccggac tcgacacett cctcttcacc tcggagtccg tgaacgaggg 120
acaccctgac aagctctgag accaggtctc agatgctgtt ctggacgctt gccttgctga 180
ggaccctgac agcaagggtg cttgcgagac ctgcaccaag accaacaatgg tcatgg 236

<210> 307
<211> 310
<212> nucleic acid
<213> Zea mays

<400> 307

cccagcacga cgagacagtc accaacgacg agatttctgg gtagagatga ggactgtgtg 60
cacacggatg ggaaccatgg cgccaccctc atctctaccc agcaccacga gacagtcacc 120
aacgacgaga ttgctgctga cctgaaggag cagctcatca agccagtcac ccccgagcag 180
tacctcgacg agaagacaat cttacacctc aaccctctct gccgcttcgt catcgccgga 240

cctcacggcg aggtgggcnt nactggccgg angatntcat cganannagg tgtttgggga 300
nccacggggg 310

<210> 308
<211> 185
<212> nucleic acid
<213> Zea mays
<400> 308

cctgacagca aggttgcttg cgagacctgc accaagacca acatgggtcat ggtctttggt 60
gagatcacca ccaaggccaa tgtcgactac gagaagattg tcagggagac atgccgcaac 120
attggtttcg tgtcgaacga tgtcgggctt gacgctgacc actgcaagtg cttgtgaaca 180
ttgag 185

<210> 309
<211> 272
<212> nucleic acid
<213> Zea mays
<400> 309

cogtttgctt cttctccctc ttgccgggtcc cgaataaaga gcagcagcgc aagaggtcgg 60
tagagcgaga agaaggcaat ggccggccgag agcttccttt tcacctcggg agtccgtgaa 120
cgagggggcac cccgacaagc tgtgcgacca ggtgtcggac gccgtgcttg acgcatgcct 180
cgcgaggagc cccgacagca aggtggcctg cgagacctgc accaagacca acatgggtgat 240
ggtgttcggc gagatcacga ccaaggcgac cg 272

<210> 310
<211> 231
<212> nucleic acid
<213> Zea mays
<400> 310

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gaagaaggca atggcgggccg agagcttcct tttcacctcg gagtccgtga acgaggggca 120
ccccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc tcgcgaggga 180
ccccgacagc aaggtggcct gcgagacctg caccaagacc aacatgggtga t 231

ctgaggaccc tgacagcaag gttgcttgag agacctgcac caagacaaca tggatcatggt 300
ctttg 305

<210> 314
<211> 237
<212> nucleic acid
<213> Zea mays
<400> 314

gaagattntc atggatacct gccgcancat tngntttgtg atcanacgat gctgggnttg 60
acgccnacca ctgnaagggtg cncgtgaana ttgatcanca gtncnctgat attgctcang 120
gtntncatgg ccacttcacc aagcgccccg aggagattgg agctggngac cagggacaca 180
tgttcgggta tgctaccnan tagaccntg agttgatgcc cctcagccat gtccttg 237

<210> 315
<211> 280
<212> nucleic acid
<213> Zea mays
<400> 315

tctcatocca ctgactccgn ccaccacccc gcccgctgcc tccgcggat ctgctcggac 60
tcggatccgc ccgaccacga ccaccccgcg ccgcccgcgc gcagagcagc agatcagaga 120
agatggccgg actcgacacc ttctcttca cctcggagtc cgtgaacgag ggacaccctg 180
acaagctctg cgaccaggtc tcagatgctg ttctggacgc ttgccttgct gaggaccctg 240
acagcaagggt tgcttgcgag acctgcacca agaccaacat 280

<210> 316
<211> 269
<212> nucleic acid
<213> Zea mays
<400> 316

cgtgaaggag aacttcgact tcaggccccg gatgatcagc atcaacctcg acctgaagaa 60
gggcggcaac aggttcatca agaccgccgc ctacggccac ttcgccgtg acgacgccga 120
cttcacctgg gaggtggtga agcccctcaa gttcgacaag gcatcggctt aagggtggga 180
gtgtcactgt ggacatgagg actaccttcc tctggctctg ctgttacctg caagcattgc 240

269

<400> 317

<400> 318

<400> 319

114

<210> 323
 <211> 280
 <212> nucleic acid
 <213> Zea mays

 <400> 323

 ggggagccca tgggtggtggc gctttctccg gncaccgtag gtgtncaatg atgatcttgc 60
 ggccagtgag gccatcattg acacctacgg tggctgggga gcccatngtg gtggcgcttt 120
 ctccggcaag gacccaacca aggttgaccg cagcggagcc tatgtcgcaa ggctggctgc 180
 caagagcatc gttcgccagc ggccttgctn cgccgcgcca tcgtccaggt gtcttacgcc 240
 atcggntggc ccgagcctct ctccgtgttc gtcgacacta 280

<210> 324
 <211> 273
 <212> nucleic acid
 <213> Zea mays

 <400> 324

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 gagecgccag cttgccccag gttggtagag cgagcgagaa gaaggcaatg gcggcggaga 120
 gcttctgtt cacctcggag tccgtgaacg aggggcaccc agacaagctg tgcgaccagg 180
 tgtcggacgc ggtgctggac gectgcctgg cgcaggaccc cgacagcaag tggctgcgag 240
 acctgcacca agacgaacat ggtgatggtg ttc 273

<210> 325
 <211> 215
 <212> nucleic acid
 <213> Zea mays

 <400> 325

 ctccctcttg ccggtcccga ataaagagca gcagcgcaag aggtcggtag agcgagaaga 60
 aggcaatggc ggccgagagc ttccttttca cctcggagtc cgtgaacgag gggcaccocg 120
 acaagctgtg cgaccaggtg tcggacgccg tgcttgacgc atgcctcgcg caggaccccg 180
 acagcaaggt ggcctgcgag acctgcacca agacc 215

<210> 326

<211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 326

 gggagaactt cgacttcagg cccgggatga tcagcatcaa cctcgacctg aagaagggcg 60
 gcaacagggt catcaagacc gccgcctacg gccacttcgg ccgtgacgac gccgacttca 120
 cctgggaggt ggtgaagccc ctcaagttcg acaaggcatc ggcttaaggt tgggagtgtc 180
 actgtggaca tgaggactac ctctctctgg ctctgtgttt acctgcaagc attgctgctg 240
 ctggatgtgt gtgtttgatc agtgactggc tgctgtcca tagaagatga a 291

<210> 327
 <211> 173
 <212> nucleic acid
 <213> Zea mays

 <400> 327

 ggtgagatca ccaccaaggc caatgtcgac tacgagaaga ttgtcaggga gacatgccgc 60
 aacattggtt tcgtgtcgaa cgatgtcggg cttgacgctg accactgcaa gtgcttgtga 120
 acattgagca gcagtcacct gatattgtc aggggtgtga cggccacttc acc 173

<210> 328
 <211> 156
 <212> nucleic acid
 <213> Zea mays

 <400> 328

 angaggggca ccccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc 60
 tcgcgagga ccccgacagc aaggtggcct gcgagacctg caccaagacc aacatgggtga 120
 tgggtttcgg cnagatcacg accanggcga ccgtnt 156

<210> 329
 <211> 178
 <212> nucleic acid
 <213> Zea mays

 <400> 329

 cccacggcgg gggcgcttc tccggcaagg acccgaccaa ggtggaccgc agcggggcct 60

acgtcgcgag gnaggctgcc aagagcatcg tcgccgccgg cctcgcccgc ngcgccattg 120

tccaggtctc ctacgccatc ggctgccccg ancccccttc ggtgttcgtg gacacgta 178

<210> 330

<211> 176

<212> nucleic acid

<213> Zea mays

<400> 330

tagttctaga tcggcaagng cnnccnttgn canatgttcg ggtacgccac cgacgagacc 60

cccagagctga tgccgctgag ccacgtgctg gccaccaagc tgggcgcgcg cctcaccgag 120

gtgcgcaaga acggcacctg cgcttggtg aggcccgacg gcaagaccca ggtgac 176

<210> 331

<211> 263

<212> nucleic acid

<213> Zea mays

<400> 331

cttctcnctc ttgccggttn cgnntaaaga ncagcagcgc aagaggctcg tagagcgatg 60

aagaaggcaa tggcggccgn nagcttcctt ttcacctcng antccgtgaa cgaangggcan 120

ccnganaagc tgtgcncca ngntcggac gccgtgcttg acgcatgcct cgcgcaggan 180

cccacagca aggtggatgc gagacctgca taagaccaac atggtgatgg tgttcgncga 240

gatcacgacc aaggcgnccg tgg 263

<210> 332

<211> 225

<212> nucleic acid

<213> Zea mays

<400> 332

cgtttgcctc ttctccctct tgccggtccc gaataaagag cagcagcgca agaggctcgt 60

agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120

aggggcaccc cgacaagctg tgcgaccagg tgcggaogc cgtgcntgac gcatgcctcg 180

cgcaggaccc cgacagcaag gtggcctgcg agacctgcac canga 225

<210> 333

<211> 331
 <212> nucleic acid
 <213> Zea mays

 <400> 333

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 catctcatcc cactgactcc ncccacccac ccgcccgcng cctccgcggg atctcgtcgg 120
 actcggatcc gncgaccac gaccaccccg cgtcgcggcc gcgcanagca gcagatcaga 180
 gaagatggcc ggactcgaca ccttcctctt cacctcggag tccgtgaacg agggacaccc 240
 tgacaagctc tgcgaccagg tctcagatgc tgttctggac gcttgccctg ctgaggaccc 300
 tgacagcaag gttgcttgcg agacctgcac c 331

<210> 334
 <211> 166
 <212> nucleic acid
 <213> Zea mays

 <400> 334

 ggaagaccca ggtgacagtg gagtaccgca acgaggggtg cgccatgggtt cccatccgtg 60
 tgcacacagt cntcatctct acccagcacg acgagacagt caccaacgac gagattgctg 120
 ctgacctgaa ggagcacgctc atcaagccag tcatccccga gcagta 166

<210> 335
 <211> 170
 <212> nucleic acid
 <213> Zea mays

 <400> 335

 ccattggttcc catccgtgtg cacacagtcc tcattctctac ccagcacgac gagacagtca 60
 ccaacgacga gattgctgct gacctgaagg agcacgtcat caagccagtc atccccgagc 120
 agtacctcga cgagaagaca atnttcacc tcaaccngtt ggnggttcgt 170

<210> 336
 <211> 247
 <212> nucleic acid
 <213> Zea mays

 <400> 336

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gggcactttg ggagggacga ccccgacttc acctgggagg tggatgaagcc cctcaaggcg 120
gagaagccgt cttctgcatg aggcgcctcc tctgtttcgg aagaagcttt tggctctggc 180
tgcttgcgct ctatcatgct tttttatggc tcctacgtgt tgtgattctt gatctgcccc 240
ttgctta 247

<210> 337
<211> 196
<212> nucleic acid
<213> Zea mays

<400> 337

gcagtnccct gatatngctc aggggtgtgca tggccacttc accaagcgcc ccgaggagat 60
tngagcnggt gaccagggac acatgttang gtatgcgacc gatgagaacnc ctgagttgat 120
gcccctcagc catgtccttg ccaccaagct aggtgcacgt ntcaaccgag gtccgcaaga 180
acggaaccng ccactg 196

<210> 338
<211> 211
<212> nucleic acid
<213> Zea mays

<400> 338

ctccctcttg ccgggtccga ataaagagca gcagcgcaag aggtcggtag agcgagaaga 60
aggcaatggc ggccgagagc ttctttttca cctcggagtc cgtgaacgag gggcaccocg 120
acaagctgtg cgaccaggtg tcggacgccg tgcttgacgc atgcntcgcg caggaccocg 180
acagcaaggt ggcttgcgag acctncacca a 211

<210> 339
<211> 302
<212> nucleic acid
<213> Zea mays

<400> 339

gacgagacga gttaccatct catcccaact ccggaacgaa caagttacca tctcatccca 60
actnccgcct tcgaccggat ctncgtcggc ctcggatccg cccgaccacc ccgcgccgcc 120

gcagatcaaa gaagatggca gctgtcgaca cattcctctt cacctcggag tctgtgaacg 180
 agggacaccc tgacaagctc tgtgaccagg tctcagatgc cgttcttgac gcttgccttg 240
 ctgaggaccc tgacagcaag ttgcttgtga gactgcacca agaccaacat ggtcatggtc 300
 tt 302

<210> 340
 <211> 263
 <212> nucleic acid
 <213> Zea mays
 <400> 340

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 tcatcaagac cgccgcctac ggccacttcg gccgtgacga cgccgacttc acctgggagg 120
 tgggtgaagcc cctcaagttc gacaaggcat cggcttaagg ttgggagtgt cactgtggac 180
 atgaggacta ccttcctctg gctctgctgt tacctgcaag cattgctgct ctggatgtgn 240
 gtgttaatca ganactgctg etc 263

<210> 341
 <211> 300
 <212> nucleic acid
 <213> Zea mays
 <400> 341

cttcgnactt caggcccggg atgatcagca tcaacctcga cctgaagaag ggccggcaaca 60
 ggttcatcaa gaccgccgcc tacggccact tcggccgtga cgacgccgac ttcacctggg 120
 aggtggtgaa gccctcaag ttcgacaagg catcggtta aggttgggag tgtcactgtg 180
 gacatgagga ctaccttct ctggctctgc tgttacctgc aagcattgct gctgctggat 240
 gtgtgtgttt gatcagtgc tggtgctgc tccatagaag atgaacggag agaaggatga 300

<210> 342
 <211> 249
 <212> nucleic acid
 <213> Zea mays
 <400> 342

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ggtanagcga gaagaaggca atggcggccg agagttcctt ttcacntcgg agtccgtgaa 120
cgagggggcac cccgacaagc tgtgcgacca gntgtcggac gccgtgcttg angcatgnct 180
cgcgaggac cccgacagca aggtggcctg cgagactgca ccaagaccaa nntggatgatg 240
gtgttcggc 249

<210> 343
<211> 196
<212> nucleic acid
<213> Zea mays
<400> 343

caatctncca cntcaaccg nctgagncgc ttcgncatcg gcggaacctca cggcgacgcn 60
ggnetcantg gccggaagat acatcatcga cacctacggn ggctggggag cccacggcgg 120
gggagcctnc tccggcaagg ncccgaccaa ggtggacngc agcggggcct acgtcgcgan 180
gnaggctgcc aagagc 196

<210> 344
<211> 249
<212> nucleic acid
<213> Zea mays
<400> 344

atcaacctcg acctcaagaa aggcggcaac gggcgctacc tcaagacggc ggctacggg 60
cactttggga gggacgaccc cgacttcacc tgggaggtgg tgaagcccct caaggcggag 120
aagccgtctt ctgcagaggc gcctcctctg ttttgaaga agcttttggc ctgncctggt 180
ctggtctggt gtgccncgcg ctctatcatg cttttttatg gctcctactt gtgattcttn 240
atctgcccc 249

<210> 345
<211> 143
<212> nucleic acid
<213> Zea mays
<400> 345

cgactcctac ggcaccggga ccatccccga caaggagatc ctaaagatcg tcaaggagaa 60
cttcgacttc aggccaggga tggtcacat caacctcgac ctcaagaagg gcggcaacag 120

gttcatcaag accgccgcat ang

143

<210> 346
<211> 142
<212> nucleic acid
<213> Zea mays

<400> 346

ccgagcctct ctccgtgttc gtngacacgt acggcaccgg cgcgatcccc gacaaggaga 60

tcctcaagat tgtcaaggag aacttcgatt tcaggcctgg catgatcatc atcaaccttg 120

acctcaagaa aggcggcaac gg 142

<210> 347
<211> 288
<212> nucleic acid
<213> Zea mays

<400> 347

cgagacgagt naccatntca tncccaacnc cggaacgaac aagttaccat ntcatnccca 60

acncccgctt cgaccggatc tccgtcggac tccggatccg cccgaccacn ccgcgccgcc 120

gcagatcaaa gaagatggca gctgtcgaca cattcctctt cacntcggag tctgtgaacg 180

agggacacnc tgacaagctc tgtgaanagg tctcagatgc cgttcttgac gcttgcnttg 240

ctgaggaccc tgacagcaag gttgcttggt agaccngcac caagacca 288

<210> 348
<211> 288
<212> nucleic acid
<213> Zea mays

<400> 348

gagacgagac gagtcccctc cccccacctc gcctcaccca accggaacga acaagttaca 60

atctcatccc aaccccgctt ccgaccggat ctgctcggac tcggatccgc ccgaccaccc 120

cgcgccgccg cagatcaaag aagatggcag ctgtcgacac attcctcttc acctcggagt 180

ctgtgaacga gggacaccct gacaagctct gtgaccaggt ctcagatgcc gttcttgacg 240

cttgcnttgc tgaggaccct gacagcaagg ttgcttggtga gacctgca 288

<210> 349

<211> 147
 <212> nucleic acid
 <213> Zea mays

 <400> 349

 ccgacaagga gatcctaaag atcgtcaagg agaacttcga cttcaggcca gggatgatca 60
 ccatcaacct cgacctcaag aagggcggca acaggttcat caagaccgcc gcatacggcc 120
 actttggccg tgacgacgcc gacttca 147

<210> 350
 <211> 264
 <212> nucleic acid
 <213> Zea mays

 <400> 350

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 agaaggcaat ggccggccgag agcttccctt tcacctcgga gtccgtgaac gaggggcacc 120
 ccgacaagct gtgacgaccag tgtggacgcc gtgcttnacg catgcctcgc gcagaccccg 180
 acagcaaggt ggcctgacgag acctgcacca agaccaacaa tgtgatngtg ttcggcgaga 240
 tnagancaaa gngactgnga tnca 264

<210> 351
 <211> 235
 <212> nucleic acid
 <213> Zea mays

 <400> 351

 cgctctttct cctccctcct gccgggtcct tantaaagag cagcagcgca agaggttggt 60
 agagcgagcg agaagaaggc aatggcggcg gagagcttcc tgttcacctc ggagtccgtg 120
 aacgaggggc acccagacaa gctgtgcgac caggtgtcgg acgcggtcnt gacggcttcc 180
 tgngncagga ccccgacagc aaggtggcct gcgagacctg caccaagang aacat 235

<210> 352
 <211> 211
 <212> nucleic acid
 <213> Zea mays

 <400> 352

cgtttgccctc ttctccctct tgccgggtccc gaataaagag cagcagcgca agagggtcgg 60
agagcgagaa gaaggcaatg gcggccgaga gcttcctttt cacctcggag tccgtgaacg 120
aggggcaccc cgacaagctg tgcgaccagg gtgcggacgc cgtgcttgac gcatgcntcg 180
cgcaggaccc cgacagcaag gtggcctgcn a 211

<210> 353
<211> 212
<212> nucleic acid
<213> Zea mays
<400> 353

gtttgcntct tctccctctt gccgggtccc aataaagagc agcagcgcaa gaggtcggta 60
gagcgagaag aaggcaatgg cggccgagag cttccttttc acatcggagt ccggtgaacga 120
gggncaaccc gacaagctgt gcgaccaggt gtgcggacgc gtgcttgacg catgcntcgc 180
gcaggacccc gacagcaagg tggcctgcga ga 212

<210> 354
<211> 490
<212> nucleic acid
<213> Zea mays
<400> 354

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tcaagaaggg cggcaacagg ttcatacaaga ccgccgcata cggccacttt ggccgtgacg 120
acnccgactt cacctgggag gtgggtcaagc ccctaaagaa ggcataccgct taagaatgta 180
ttgggaagtt cactggacat gaggttcac ttcgtctggc tctgctgata cctgcaagga 240
tnnnnnnnnn nnnnnnnnnc ctggatgtgt gtttgatcag tgactggctg ctctgctcca 300
tagaagatga atgaagagag agatggtgaa naaggctttg gcaaattggca attgccgaac 360
aagccatgtc gcnccactga ccggcttaat gattggtata atttggtgtg gcaacancca 420
ggattaatgc ccctggncctt ttatcnttac tactaanttg ggctngtccg gtatctaatt 480
ttctttccct 490

<210> 355
<211> 389
<212> nucleic acid

<213> Zea mays

<400> 355

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accacgcgt ccgcggaagc gtgggcctcg acctcaagaa aggcggcaac gggcgctacc 120

tcaagacggg ggcctacggg cactttggga gggacgaccc cgacttcacc tgggaggtgg 180

tgaagccct caaggcgag aagccgtctt ctgcatgagg cgctcctct gtttcggaag 240

aagcttttgg tctggtctgc ctgctctta tcatgcttt ttatggtcc tacgtgttgt 300

gattcttgat ctgccccttg cttatcattt gtattgtact gtcactgtcc taataagtgg 360

tacgtgtgag gggctgtatt gtgtctgct 389

<210> 356

<211> 289

<212> nucleic acid

<213> Zea mays

<400> 356

gtcggatctg agacgagacg agtcccctcc cccacctcg cctcacccaa ccggaacgaa 60

caagttacaa tctcatccca acccgcctc gaccggatct cgtcggactc ggatccgccc 120

gaccaccccg cgcgcgcga gatcaaagaa gatggcagct gtcgacacat tctctttcac 180

ctcggagtct gtgaacgagg gacaccctga caagctctgt gaccaggtct cagatgccgt 240

tcttgacgct tgccttgctg aggaccctga cagcaaggtt gcttgtag 289

<210> 357

<211> 264

<212> nucleic acid

<213> Zea mays

<400> 357

gggatgatca gcatcaacct cgacctgaag aagggcgga acaggttcat caagaccgcc 60

gcctacggcc acttcggccg tgacgacgcc gacttcacct gggaggtggg gaagcccctc 120

aagttcgaca aggcacggc ttaaggttg gagtgctact gtggacatga ggactacctt 180

cctctggctc tgctgttacc tgcaagcatt gctgctgctg gatgtgtgtg tttgatcagt 240

gactggctgc tgctccatag aaga 264

<210> 358
 <211> 263
 <212> nucleic acid
 <213> Zea mays

 <400> 358

 gggatgatca gcatcaacct cgacctgaag aagggcggca acaggttcat caagaccgcc 60
 gcctacggcc acttcggccg tgacgacgcc gacttcacct gggagggtgt gaagcccctc 120
 aagttcgaca aggcacgcgc ttaagggttg gagtgtcact gtggacatga ggactacctt 180
 cctctggctc tgctgtgacc tgcaagcctt gctgctgctg gatgtntgtg tttgatcagt 240
 gantggctgc tgctccatag ang 263

<210> 359
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 359

 gagacgagac gagttaccat ctcatcccaa ctccggaacg aacaagttac catctcatcc 60
 caactccgcc tcgaccggat ctctcggac tcggatccgc ccgaccaccc cgcgccgccg 120
 cagatcaaag aagatggcag ctgtcgacac attcctcttc acctcggagt ctgtgaacga 180
 gggacaccct gacaagctct gtgaccaggt ctcatatgcc gttcttgacg cttgccttgc 240
 tgaggaccct gacagcaagg ttgcttgc 268

<210> 360
 <211> 289
 <212> nucleic acid
 <213> Zea mays

 <400> 360

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 acaagttaca attctcatcc caaccccgcc tcgancggat ctctcggac tcggatccgc 120
 ccgaccaccc cgcgccgccg cagatcaaag aagatggcag ctgtcgacac attcctcttc 180
 acctcggagt ctgtgaacga gggacaccct gacaagctct gtgaccaggt ctcatatgcc 240
 gttcttgacg cttgccttgc tgaggaccct gacagcaagg ttgcttgc 289

<210> 361
 <211> 252
 <212> nucleic acid
 <213> Zea mays
 <400> 361
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 gaagaaggca atggcggccg agagcttcct tttcacctcg gagtccgtga acgaggggca 120
 cccgacaagc tgtgcgacca ggtgtcggac gccgtgcttn acgcatgcct cggcaggacc 180
 ccgaaacaag tggctggaga ccttcaccaa gancaatntg gtgatggtgt tcggngaate 240
 acgacaaggc ga 252

<210> 362
 <211> 246
 <212> nucleic acid
 <213> Zea mays
 <400> 362
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 cggccacttc ggccgtgacg acgccgactt cacctgggag gtggtgaagc ccctcaagtt 120
 cgacaaggca tcggcttaag gttgggagtg tcaactgtga catgaggact accttcctct 180
 ggcctctgctg ttacctgcaa gcattgctgc tgctggatgt gtgtgtttga tcagtgactg 240
 gctgct 246

<210> 363
 <211> 240
 <212> nucleic acid
 <213> Zea mays
 <400> 363
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 ttgtccagga tntcctacgc cangcggctg cccanancec ctttcggtgt tcgtggacac 120
 gtacggcacc ggcgcgatcc ccgacaagga gatcctgaag atcgtgaagg agaacttcga 180
 cttcaggccc ggcattgntca tcataacact cgacctcaag aaaggcngca agggcgtact 240

<210> 364

<211> 270
 <212> nucleic acid
 <213> Zea mays

 <400> 364

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 tctcatccca accccgcctn cgaccggatc tctcggaact cggatccgcc cgaccacccc 120
 gcgcgcgcgc agatcaaaga agatggcagc tctcgacaca ttctcttca cctcggagtc 180
 tgtgaacgag ggacaccctg acaagctctg tgaccaggtc tcagatgccg ttcttgacgc 240
 ttgccttgct gaggaccctg acagcaaggt 270

<210> 365
 <211> 252
 <212> nucleic acid
 <213> Zea mays

 <400> 365

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 ctctgcccg gtcttaata aagagcagca gcgcaagagg ttggtagagc gagcgagaag 120
 aaggcaatgg cggcggagag cttctgttc acctcggagt ccgtgaacga ggggcaccca 180
 gacaagctgt gcgaccaggt gtcngacgcn gtgctggacg cctgcctggc gcaggacccc 240
 gacagcaagg tg 252

<210> 366
 <211> 320
 <212> nucleic acid
 <213> Zea mays

 <400> 366

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 tcatcccaact gactcngcc acccactann ncnntgcctc cgccggatct cgtcggactc 120
 ggatccgccc gaccacgacc accccgcgcc gccgccgcgc agagcagcag atcagagaag 180
 atggccggac tcgacacctt cctcttnacc tcggagtccg tgaacgaggg acaccctgac 240
 aagctctgcg accaggtctc agatgctggt ctggacgctt gccttgctga ggaccctgac 300
 agaaagttgt tgcgagactg 320

<210> 367
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 367

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 gttaccatct catcccaact cgcctcgac cggatctcgt cggactcgga tccgcccgcac 120
 cccccgcgc cgcgcgagat caaagaagat ggcagctgtc gacacattcc tcttcacctc 180
 ggagtctgtg aacgagggac accctgacaa gctctgtgac caggtctcag atgccgttct 240
 tgacgcttgc cttgctgagg accctgacag caag 274

<210> 368
 <211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 368

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 atctcatccc aactccgctt cgaccggatc tcgtcggact cggatccgcc cgaccacccc 120
 gcgcgcgcgc agatcaaaga agatggcagc tgtcgacaca ttcctcttca cctcggagtc 180
 tgtgaacgag ggacaccctg acaagctctg tgaccaggtc tcagatgccg ttcttgacgc 240
 ttgccttgct gaggaccctg acagcaagtt g 271

<210> 369
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 369

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 atactgcac ccaacccgc ctcgaccgga tctcgtcgga ctcggatccg cccgaccacc 120
 ccgcgcgcgc gcagatcaaa gaagatggca gctgtcgaca cattcctctt cacctcggag 180
 tctgtgaacg agggacaccc tgacaagctc tgtgaccagg tctcagatgc cgctcttgac 240
 gcttgccctg ctgaggaacc tgacagcaag gttg 274

<210> 370
 <211> 203
 <212> nucleic acid
 <213> Zea mays

 <400> 370

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 gaagaaggca atggcggcgc agagcttctt tttcaentcg ggtccgtga acgaggggca 120
 cgccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc tcgcgagga 180
 cnccgacagc aagtggcctg cga 203

<210> 371
 <211> 201
 <212> nucleic acid
 <213> Zea mays

 <400> 371

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 gagcgagcga gaagaaggca atggcggcgc agagcttctt gttcacctcg ggtccgtga 120
 acgaggggca cccagacaag ctgtgcgacc aggtgtcgga cgcggtgctg gacgcctgcc 180
 tggcgagga ccccgacagc a 201

<210> 372
 <211> 307
 <212> nucleic acid
 <213> Zea mays

 <400> 372

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 gccacttcgg ccgtgacgac gccgacttca cctgggaggt ggtgaagccc ctcaagttcg 120
 acaaggcatc ggcttaaggt tgggagtgtc actgtggaca tgaggactac ctctctctgg 180
 ctctgctgtt acctgcaagc attgctgctg ctggatgtgt gtgtttgatc agtgactggc 240
 tgctgctcca tagaagatga acggagagaa ggatgatgaa gggctttggc aatcgccgct 300
 gcaactg 307

<210> 373

<211> 283
 <212> nucleic acid
 <213> Zea mays

 <400> 373

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 gogatccccg acaaggagat nctcaagatt gtcaagngaa ttcgatttca ggcttggcat 120
 gatcatcatc aaccttgact caagaaaggc ggcaacgggc gctactcaag acgcggccta 180
 cggcactttg gaaggagacc tgattcactg ggagtgggga accatcaatg gagaacttgc 240
 tennngctat tatataantt gtgcgttctg actatngtta nag 283

<210> 374
 <211> 181
 <212> nucleic acid
 <213> Zea mays

 <400> 374

 cttagccggtc ccgaataaaag agcanncagc gcaagagntc ggtagagcga caagaaggca 60
 atggcgggcg agagcttctt ttacacctcg gagtccgtga acgaggggca ccccgacaag 120
 ctgtgcgacc aggtgtcgga cgccgtgctt gacgcagtcg tcgcgcagga ccccgacagc 180
 a 181

<210> 375
 <211> 201
 <212> nucleic acid
 <213> Zea mays

 <400> 375

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 gtagagcgag cgagaagaag gcaatggcgg cggagagctt cctgttcacc tcggagtccg 120
 tgaacgaggg gcacccagac aagctgtgcg accaggtgtc ggacgcgggtg ctggacgcct 180
 gcctggcgca ggaccccgac a 201

<210> 376
 <211> 216
 <212> nucleic acid
 <213> Zea mays

<400> 376

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gcgcaagagg ttggtagagc gagcgagaag aaggcaatgg cggcggagag cttcctgttc 120
acctcggagt ccgtgaacga gtggcaccca gacaagctgt gcgaccaggt gtcggacgcg 180
gtgctggacg cctgcctggc gcaggacccc gacagc 216

<210> 377

<211> 130

<212> nucleic acid

<213> Zea mays

<400> 377

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gtccccgat attgcnacgg gtgtgcacgg ccacttcacc aagcgccccg aggagattng 120
agctggtgac 130

<210> 378

<211> 306

<212> nucleic acid

<213> Zea mays

<400> 378

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actcggatcc gcccgaccac gaccaccccg cgtcgccgc gcgcanagca gcagatcaga 180
gaagatggcc ggactcgaca ctttctctt cacctcggag tccgtgaacg agggacaccc 240
tgacaagctc tgcgaccagg tctcagatgc tgttctggag gnttgccttg ctgaggaccc 300
tgacag 306

<210> 379

<211> 313

<212> nucleic acid

<213> Zea mays

<400> 379

gagacgagac gannnnccct ccctcaacc ggaacttggt ttaccccatc tcatccact 60

gactccan cn acccaccgc ncgtgcctc cgccggatct cgtcggactc ggatccgccc 120
gaccacgacc accccgcgtc gccgccgcgc agagcagcag atcagagaag atggccggac 180
tcgacacctt cctcttcacc tcggagtccg tgaacgaggg acaccctgac aagctctgga 240
ccaggtctca gatgctgttc tggacgcttg ccttgctgag gacctgacag caaggttgct 300
tgggagacct gca 313

<210> 380
<211> 134
<212> nucleic acid
<213> Zea mays
<400> 380

gengacctca nggagcacgt catcaagcnc gtgatccctg agaagtacct cgacgagang 60
accatcttcc acctcaacct gtccggggcgc ttctgtcatcg gcngggcccca cggtnacncc 120
ngentcacng gtgc 134

<210> 381
<211> 294
<212> nucleic acid
<213> Zea mays
<400> 381

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taccatctca tcccaactcc gctctgaccg gatctctgtcg gactcggatc cggccgacca 120
ccccgcgccg ccgcagatca aagaagatgg cagctgtcac acattcctct tcacctcgga 180
gtctgtgaac gagggacacc ctgacaagct ctgtgaccag gtctcagatg ccgttcttga 240
cgcttgcttt gctgaggacc ctgacagcaa ggttgcttgt gagactgcac caag 294

<210> 382
<211> 164
<212> nucleic acid
<213> Zea mays
<400> 382

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gtccgcaaga gcggaaacct gccctggct acaggcctga tgggaagacc caggtgacag 120

tcgagtaccg cantgagggt ggtgccatgg tccccatccg tntc 164

<210>	383
<211>	247
<212>	nucleic acid
<213>	Zea mays

<400> 383

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nggnnnaggt	tgatnatgat	catgccaggc	ctgaaatoga	agttctcctt	gacaatcttg	120
aggatctcct	tgtcggngat	cagngccagt	gcacgtacag	ngtcnacgaa	cacnganaga	180
ngctcgggca	ctccnntnnc	ngtaagacan	ctggacnatg	gttagtgnaa	gtnatncnt	240
tgaanac						247

<210>	384
<211>	207
<212>	nucleic acid
<213>	Zea mays

<400>	384
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cagcagcgcga agaggtcggg agagcgagaa gaaggcaatg gcggccgaga gcttcctttt 120
cacctcggag tccgtgaacg aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc 180
cgtgcttgac gcatgcctcg cgcagga 207

<210>	385
<211>	292
<212>	nucleic acid
<213>	Zea mays

<400> 385

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cccatctcat	cccacagnct	ccacccannc	gcccgctgcc	tccgcgggat	ctcgtcggac	120
tcggatccgc	ccgaccaccc	cgcgcgcgcg	ccgcgcagag	cagcagcaga	tcagagaaga	180
tggccggact	cgacaccttc	ctcttcacct	cggagtcogt	gaacgagggg	caccttgaca	240
agctctgcga	ccagggtctca	gatgctgttc	tggacgcttg	ccttgctgag	ga	292

<210> 386
 <211> 142
 <212> nucleic acid
 <213> Zea mays

 <400> 386

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 aagacccagg tgacgggtgga gtacntgaac gagggcgggc ccatgggtgcc cgtccgcntg 120
 cacaccgtgt catctccaca ca 142

<210> 387
 <211> 137
 <212> nucleic acid
 <213> Zea mays

 <400> 387

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 ctgagccacg ngntggccac caagntgggc gcgcgcntca ccgangngcg caagaacggc 120
 acntggcnen tggngga 137

<210> 388
 <211> 159
 <212> nucleic acid
 <213> Zea mays

 <400> 388

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 ccccgacttc acctggnagg tggatgaagc cctcaaggcg gagaagccgt cttctgcatg 120
 aggcgccctc tctnttttgn aangancttt tggtcnggt 159

<210> 389
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 389

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 ctcatcccaa ccccgccctcg accggatctc gtcggactcg gatccgccc accaccccgc 120

gccgccgcag atcaaagaag aggcagctgt cgacacattc ctcttcacct cggagctctgt 180
gaacncggga caccctgaca agctctgtga ccangtctca gatgccgttc ttgacgcttg 240
ccttgctgag gaccctgaca gcaaggtt 268

<210> 390
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 390

aagaagggcg gcaacagggt catcaagacc gccgcctacg gccacttcgg ccgtgacgac 60
gccgacttca cctgggaggt ggtgaagccc ctcaagttcg acaaggcatc ggcttaaggt 120
tgaggagtgtc actgtggaca tgaggactac ctctctctgg ctctgctgtt acctgcaagc 180
attgctgctg ctggatgtgt gtgtttgatc agtgactggc tgctgcttcc atagaagatg 240
aaggagagaa ggatgatgaa gggctttggc aatcgccgcg ca 282

<210> 391
<211> 272
<212> nucleic acid
<213> Zea mays

<400> 391

caacggggcg tacctcaaga cggcggccta cggccacttt ggaagggacg accctgactt 60
cacctgggag gtggtgaagc cactcaagtc ggagaaacct tctgcctaag gcggcctttt 120
tttttcagta agaagctttt ggtggtctgc tgtgcttaat catgctttta tatggcttct 180
acatgtttgt gttctttctt gatctgcacc gcgcttatcg tttgtgttgt actgccctaa 240
taagtgggtgc tatgaggact gtttctgggt tt 272

<210> 392
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 392

cggatctgag acgagacgag acgannnncc ctcccctcaa ccggaacttg ttttaccoca 60
tctcatocca ctgnoctccn gccaccacc cgcncgctgc ctccgccgga tctcgtcgga 120

ctcggatccg ccgaccacg accaccccg gtcgccgcg cgcagagcag cagatcagag 180
aagatggccg gactcgacac cttcctcttc acctcggagt ccgtgaacga gggacaccct 240
gacaagctct ggcaccaggt ctcagatgct gttctggacg cttgccttgc t 291

<210> 393
<211> 531
<212> nucleic acid
<213> Zea mays

<400> 393

agnnnnnnn natntaatga atttnangaa tgctctacch gnaattcccg ggtcgaccca 60
cgcgtccgnc cacnecgtec angagatcct caagatcgng aaggagaact tcgacttcag 120
gcccngnatg atcagcatca acctngacct gaanaaggnc ggcaacaggt tcatcaagac 180
cgacgcctac agtcacttcn gncgtgacga cnccgactta cctggnaggt ggtgaacccc 240
tcaagttcga caaggcatcg ncttaaggct gngaagtgc cactgtggac attgaggact 300
accttactct ggctctgntg gtacctgcaa agcattggct gctgatggat gtntgngnct 360
gatcaagnga ctggctgctg cttcatanna gatntaccg aganaaagat gatgnataaa 420
ggcttnggca atcggcggtt canctgnaac ccatgccatt ccgcttanng aatggggata 480
anttggcttg gaaanaanca tcattattat ggnetgaact ttcacttita c 531

<210> 394
<211> 572
<212> nucleic acid
<213> Zea mays

<400> 394

gggggnnnng gnaacttcta tntcgnccg cacggtccaa aaaatcccgg ggtccgaccc 60
acgcgttccg aggcnaacttt tctcccggca aagggaccca aaccaaagg tttgaaccnc 120
aagccgggaa ccctaatttt cgcaaaggg caanggctng cccaaagaac caatccgtcc 180
gcccgaagccg ggccctttgc ctccgcccgc cgccaatccg ttccaangat tgtcttaacg 240
ccaatccgng cgttnccccg aaacctctct ccgttggtcg tcgacacnta cggcaccngg 300
cgcgatcccc gacaaggaan atnctcaaga ttgtcaagga agaacttca tttcaggcct 360
gngcatgac atcatcaacc ttgacctcaa gaaangcggc aacggggcgc tacctcaaag 420

ccctganagc naaggtgctt gtganacctg

270

<210> 398
<211> 284
<212> nucleic acid
<213> Zea mays

<400> 398

catcaacctc gacctcaaga ngggcggaac caggttcacg aagaccgccg catacggcca 60
ctttggccgt gacgacgccg acttcacctg ggaggtggc aagcccctaa agaaggcatc 120
cgcttaagaa tgtattggga agttcactgg acatgaggtt catcttcgtc tggctctgct 180
gataacctga aggatnnnnn nnnnnnnnnn nnnnnnnnga tgtgtgtttg atcagtgact 240
ggctgctctg ctccatagaa gatgaatgaa gagagagatg gtga 284

<210> 399
<211> 297
<212> nucleic acid
<213> Zea mays

<400> 399

atctgagacg agacgnngnc nncctcncct caaccggaac ttgttttacc ccctctcatc 60
ccantgante nagnnannca cncgncgct gntccgncg gatctcttcg gactcggatc 120
cgcccgancc cgaccanccc gcgcgcgcgc cgcgagagc agcagatcag agaagatggc 180
cggactcgac accttctctt tcacctcgga gtccgtgaac gagggacacc ctgacaagct 240
ctgcgaccag gtctcagatg ctgttctgga ngttgcttgc tgangacctg acagcaa 297

<210> 400
<211> 279
<212> nucleic acid
<213> Zea mays

<400> 400

gtcggatctg agacgagacg agacgatnnc cctccccctc aaccggaact tgttttacct 60
catctcatcc cacngactcc ncccaccac ccgcccgtg cctccgcccg atctcgtcgg 120
actcggatcc gcccgaccac gaccaccccg cgtcgcgcgc gcgcagagca gcagatcaga 180
gaagatggcc ggactcgaca ccttctcttt cacctcggag tccgtgaacg agggacaccc 240

tgacaagctc tgcgaccagg tctcagatgc tgttctgga 279

<210> 401
<211> 307
<212> nucleic acid
<213> Zea mays

<400> 401

cggatctgag acgagacgag acgatnnncc ctccctcaa ccggaacttg ttttaccoca 60
tctcatccca ctgantctnc ccattccacc gcccgngcc tccgccgat ctctgtggac 120
tcggatccgc ccgaccacga ccaccccgcg tcgcccgcgc gcagagcagc agatcagaga 180
agatggccgg actcgacacc ttctcttca cctcggagtc cgtgaacgag ggacacctg 240
acaagctctg cgaccaggtc tcagatgctg ttctggacgt tgcttgctga ggacctgaca 300
gcaaggt 307

<210> 402
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 402

gtttgcctct tctccctctt gccggtcccg aataaagagc agcagcgcaa gaggtacggt 60
agagcgagaa gaaggcaatg gccgncgaga gcttctttt cacctcggag tccgtgaacg 120
angggcacc cgcacaagctg tgcgaccagg ttacaaaaan ccgtgcttga cgcattgcctc 180
gcgcagaccc cgacagcaag gtggcttncg agacttnac caagaccaca tggtangttt 240
tngnngntgg nncgncaaag nnaangntt tnanaaaat ntntnnancc c 291

<210> 403
<211> 386
<212> nucleic acid
<213> Zea mays

<400> 403

caagaaagnc ggcaacgggc cgctacctca agacgggggc gnacggccac tttggaaggg 60
acgacctga cttcacctgg gaggtggtga agccactcaa gtcggagaaa cttctgcct 120
aaggcggcct tttttttcag taagaagctt ttggtggtct gctgtgctta atcatgcttt 180

tatatggctt ctacatgttg tggttctntc ttgatctgca ccgngcttat cgnntnngtt 240
gtactgncct aataaatnng tgcttatgan gacttgtnn tggntnnnt antanngtn 300
naatgcttta aaacaatgan tgaattncaa gccannnttt ttttgagaag taannattat 360
tngntaannn gntnngnntn tnnngg 386

<210> 404
<211> 144
<212> nucleic acid
<213> Zea mays
<400> 404

tccgtgttcg tcgacacgta cggcaccggc gcgatccccg acaaggagat cctcaagatt 60
gtcaaggaga acttcgattt caggcctggc atncatcatc atcaaccttg acctcaagaa 120
aggcggcaac gggcgctacc tcaa 144

<210> 405
<211> 293
<212> nucleic acid
<213> Zea mays
<400> 405

agaacttcga cttcaggccc gggatgatca gcatcaacct cgacctgaag aagggcggca 60
acaggttcat caagaccgcc gcctacggcc acttcggccg tgaacgacgc cgacttcacc 120
tgggaggttg tgaagcccct caagtctgac aaggcatcgg cttaagggttg ggagtgtcac 180
tgtggacatg aggactacct tcctctggct ctgctgttac ctgcaagcat tgctgtgtgt 240
ggatgtgtgt gtttgatcag tgactggctg ctgtccatag aagatgaacg gag 293

<210> 406
<211> 175
<212> nucleic acid
<213> Zea mays
<400> 406

ggtcaccatc aacctcgacc tcaagaaggc cggcaacagg ttcacatcaaga ccgccgcata 60
cgccatttg gncgtgacga cgccgacttc acctgggagg tggtaagcc cctaaagaag 120
gcatccgctt aagaatgtat tgggaagttc actggacatg aggttcatct tcgtc 175

<210> 407
 <211> 219
 <212> nucleic acid
 <213> Zea mays

 <400> 407

 aggggtgtgca cggccacttc accaagcgcc ccgaggagat tggagctggt gaccaggggc 60
 acatgttttg gntgcgactg acgagacccc tgagtgatgc cctcagccat gtcttgccac 120
 caagctggtg tcgtctcacg gagtnccaag atggactgcc ctgntcagcc gtggaagacc 180
 agtgcagtga tacgnagagg tggcatgtcc acggtnnnc 219

<210> 408
 <211> 178
 <212> nucleic acid
 <213> Zea mays

 <400> 408

 gccagggatg atcaccatca acctcgacct caagangggc ggcaacaggt tcatcaagac 60
 cgccgcatac ggccactttg gctgaacgac gccgacttca cctgggaggt ggtcaagccc 120
 ctaaagaagg catccgctta agaatgtatt gggaagtcca ctggacatga ggttcac 178

<210> 409
 <211> 126
 <212> nucleic acid
 <213> Zea mays

 <400> 409

 gcaatggcgg cggagagctt cctgttcacc tcggagtccg tgaacgaggg gcacccagnc 60
 aagctgttcg ancaggtgtc tgangcggtc tggangcctt cctgnntcag gancccgaca 120
 ntaaag 126

<210> 410
 <211> 132
 <212> nucleic acid
 <213> Zea mays

 <400> 410

 gacctcaaga ngggcggcaa caggttcac 60

gacgacgccg acttcacctg ggaggtggc aagcccctaa agaaggcatc cgcttaagaa 120
tgtattggga ag 132

<210> 411
<211> 83
<212> nucleic acid
<213> Zea mays

<400> 411

gtcggangcg gtgctggang cctgcctggn gcagganncc ganagcaagg tggcctgcga 60
ganctgcacc aagangaaca tgg 83

<210> 412
<211> 133
<212> nucleic acid
<213> Zea mays

<400> 412

gcctcgaccg gatctcgtcg gactcggatc cgcccgacca ccccgcgccg ccgcagatca 60
aagaagatgg cagctgtcga cacattctc ttcacctcg agtctgtgaa ngaggganac 120
cctgacaagc tct 133

<210> 413
<211> 290
<212> nucleic acid
<213> Zea mays

<400> 413

tccgatctga gacgagacga gacggnnnnc cctcccctca accggaactt gttttacccc 60
atctcatccc agtgantcnt accacncanc cgcgcgngc ntccgngga tctngtcgga 120
ctcggatccg cccgaccacg accaccccg gtcgccgccg cgcagagcag cagatcagag 180
aagatggccg gactcgacac cttcctcttc acctcggagt ccgtgaacga gggacaccct 240
gacaagtctg cgaccaggtc tcagatgtgt ttggacgttg nttgctgagg 290

<210> 414
<211> 310
<212> nucleic acid
<213> Zea mays

<400> 414

aacagggttca tcaagaccgc cgcatacggc cactttggcc gtgacgacgc cgacttcacc 60
tgggaggtgg tcaagcccct aaagaaggca tccgcttaag aatgtattgg gaagttcact 120
ggacatgagg ttcattctcg tctggctctg ctgatacctg caaggatnnn nnnnnnnnnn 180
nnnnnnnnnn gatgtgtggt tgatcagtga ctggctgctc tgctccatag aagatgaatg 240
aagagagaga tgggtgaagaa ggctttggca aatggcaatt gccgcagcaa gccatgtcgg 300
cgccactgac 310

<210> 415

<211> 85

<212> nucleic acid

<213> Zea mays

<400> 415

ctcagggtgt gcatggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 60
gacacatggt cgggtatgcg accga 85

<210> 416

<211> 166

<212> nucleic acid

<213> Zea mays

<400> 416

gagcagcagc gcaaggngan ccgccagctt gccccaggtt ggtagancga gcnagaagaa 60
ggcaatnncg ggggagagtt cctgttcacn tcggagtccg tgaacgangg gcacccagac 120
aagctgtgcg accaggtntc ggacgcggtg ctggacncct gcntgg 166

<210> 417

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 417

aagacggcgg cctatggcca ctttgaagg gacgaccctg acttcacctg ggaggtggtg 60
aagccactgc aagtcggaga aaccttctgc ctaaggcggc cttttttttc agtaagaagc 120
ttttggtggt ctgctgtgct taatcatgct tttatanggc ttctacatgt tgtggttctt 180

tcttgatctg caccgcgctt atcgtttctg ttgtactgcc ctaataagtg gtgcttatga 240
ggactgtttc tggttttgct gcttatg 267

<210> 418
<211> 273
<212> nucleic acid
<213> Zea mays

<400> 418

acgaccccca cttcacctgg gaggtggtga agcccctcaa ggcggagaag ccgttctctg 60
catgaggcgc ctctctgttt ttggaagaag cttttggtct ggtctggtct ggtctggtgt 120
gootgcgcgc tatcatgctt ttttatggct cctacttctg attcttgatc tgcccccttgc 180
ttatcatttg tactgtactg tcaactgtcct aataagtggc acgtgtgcgg ggtcgtattg 240
tgtctgctta ttcacctaga ggattatttc tgg 273

<210> 419
<211> 57
<212> nucleic acid
<213> Zea mays

<400> 419

atcgtctctg acctgaagga gcatgtcatc aagcctgtca tccctgagca gtacctt 57

<210> 420
<211> 235
<212> nucleic acid
<213> Zea mays

<400> 420

gtcggatctg agacgagacg nngnnncct cccctcaacc ggaacttgtt ttaccccatc 60
tcateccact gactcngncc acccaccann ncantgcctc cgccggatct cgtcggactc 120
ggatccgccc gaccacgacc accccgcgcc gccgccgcgc acagcagcag atcagagaag 180
atggccggac tcgacacctt cctcttcacc tcggagtccg tgaacgaggg acacc 235

<210> 421
<211> 297
<212> nucleic acid
<213> Zea mays

<400> 421

gccaaagggat gatcaacaat ccaacntcga nctccaagaa ngggcggnaa caggttcac 60
aagaccgccc catacggcca ctttggccgt gaacgacgcc gacttcacct gggaggtggt 120
caagccccta aagaaggcat ccgttaagaa tgtattggga agttcactgg acatgaggtt 180
catcttcgtc tggctctgct gatacctgca aggatnnnnn nnnnnnnnnn nnnnnnnnga 240
ttgtgtttga tcagtgactg gctgctctgc tccatagaag atgaatgaag agagaga 297

<210> 422

<211> 88

<212> nucleic acid

<213> Zea mays

<400> 422

caoncaagag accgtcacca acgacgagat cgccgccgac ctcaaggagc acgtcatcaa 60
gcccgtgac cctgagaagt acctgcga 88

<210> 423

<211> 285

<212> nucleic acid

<213> Zea mays

<400> 423

ccgggtcgga tctgagacga gacgagttac catctcatcc caactccgga acgaacaagt 60
taccatctca tcccaactcc gctcgcaccg gatctcgtcg gactcggatc cgcccgacca 120
ccccgcgng ccgcagatca aagaagatgg cntcgtcgac acattcctct tcacctcgga 180
gtctgtgaac gagggacacc ctgacaagtc tgtgaccagg tctcagatgc cgttcttgag 240
cttgcnttgc tgaggacct gacagcaagg ttgttgag actgc 285

<210> 424

<211> 136

<212> nucleic acid

<213> Zea mays

<400> 424

accacgacca ccccgctcg ccgccgcga naggcagaga tcagagnaga tagccggatc 60
tcgacacnt cctcttcacc tcggagtccg tgaacgagg acaccctgac aagctctgcg 120

accaggtctc agatgc

136

<210> 425
<211> 217
<212> nucleic acid
<213> Zea mays

<400> 425

cgagacgagt nncctcccc cacctcgctt caccacaaccg gaacgaacaa gttacaatac 60
tcatacccaac cccgccttcg accggatctc gtcggactcg gatccgcccg accaccccgc 120
gccgccgcag atcaaagaag atggcagctg tcgacacatt cctcttcacc tcggagtctg 180
tgaacgaggg acacctgac aagctctgtg accaggt 217

<210> 426
<211> 231
<212> nucleic acid
<213> Zea mays

<400> 426

cggatctgag acgagacgag ttaccatctc atcccaactc cggaacgaac aagttaccat 60
ctcatoccaa ctccgcttcg accggatctc gtcggactcg gatccgcccg accaccccgc 120
gccgccgcag atcaaagaag atggcagctg tcgacacatt cctcttcacc tcggagtctg 180
tgaacgaggg acacctgaca agctctgtga ccaggctcaa tgccgttctt g 231

<210> 427
<211> 85
<212> nucleic acid
<213> Zea mays

<400> 427

agtacctcga ngagaagacc atcttcacc tcaacccgctc cgggcgcttc gtcacgggnn 60
ggntcgangg tgacgtnggc ctcat 85

<210> 428
<211> 142
<212> nucleic acid
<213> Zea mays

<400> 428

<210> 432
 <211> 240
 <212> nucleic acid
 <213> Zea mays

<400> 432

gggagaaatt cgtgagatct tggnnctgnt cagggcgtgc gagcncctgga atcatgggtt 60
 tcacacatag ctctcncntnt tngaatttna tgtactaatg gagtcnaagg gtggcaaaaa 120
 gtcnnncngt agtcgttcta tgggtgatga agcgccctt ggctacagca ttgaggncgt 180
 tcgacctgcc ggagcgtgaa gaantccagc tgcggcttac tcgaactgcg ngaatnagcg 240

<210> 433
 <211> 263
 <212> nucleic acid
 <213> Zea mays

<400> 433

cttgtgtcgg tagttttccg ttggctctgg ctgccttctt ctgcttctga gattccaact 60
 tgtcttgcgc atctctcctt ttctctcttc atctctcttt tcttgcaaca cgtcatccag 120
 tggcgatgtc ttcagcagat tcttgtgtct cttctcctgc ctccctatt ggctttgagg 180
 gctatgagaa gcgcctcgag atcacgttct ctgacgcgcc tgtctttgag gacccttggtg 240
 gtctgtgcnt gcgcgccttc tcc 263

<210> 434
 <211> 290
 <212> nucleic acid
 <213> Zea mays

<400> 434

taatgtgat ggcaacacaa cattagtctt gaagaagaat gaagctttct tcaagactaa 60
 tgctgatggc aacacaacat anttccaagg aaatgacgaa gctctctggt atctctgaaa 120
 ttatccctga gaaggagatc tgtgattttg acttcgaacc ctgcggctac tccatgaatg 180
 caatccatgg ctctgcgttc tccacgatcc atgtgacgcc tgaggatggg ttcagctacg 240
 ccagttatga ggttatgggc ttggatgcca ctgccctgtc ttacggtgac 290

<210> 435

<211> 258
 <212> nucleic acid
 <213> Zea mays

 <400> 435

 tggagcagga gctgcctgga ggcgggctcc tcgtgtacca tngcttctgt gntgctgaag 60
 acgtgtttgc tacctcgccc aaatctgttt tccacngcnt tganggcgag aacgtggaga 120
 gtgtcctctn tcgnntgagg agganancaa gnggnnaang ttctntgttg ggagnatnan 180
 acggngtgcc atggaggaga ngcggggatt ccttgatgag taatangggg ctctgggntc 240
 gattagcttc tgattgtt 258

<210> 436
 <211> 263
 <212> nucleic acid
 <213> Zea mays

 <400> 436

 cttgtcttaa aacaatcgca gtcttgcaag ttgttgctgc tgctcctacc cctgcctctg 60
 caattgggtt tgagggatat gagaagcgcc tcgagatcag cttctatgag gcacctgtct 120
 tcgtgaccc caacggaagg ggattgtgtg cactcttgcg tccctagatt tactctattt 180
 ctgaccatgc acggtgcacc gttgtctctg agctatcaaa cgaggacttt gactctatgt 240
 cttatctgag tcaagcctgt ttg 263

<210> 437
 <211> 266
 <212> nucleic acid
 <213> Zea mays

 <400> 437

 cccaganatg gagatctgtg acttcgactt cnagccctgt ggctactcca tgacatgctg 60
 ttcattggcnc tgcnattgtc gaccattcat gnganccnc aggacgggca tcagctaattg 120
 aaagcaacan ggncatgggc nttaanccgg ggctcctttc tcatatggtn anctggntaa 180
 ganggtgctg aagnnntttt gncnactga ntncnctgtt gncgtaacct atcttccggt 240
 gatcgggcaa tgcgaagacc tggggg 266

<210> 438

<211> 281
 <212> nucleic acid
 <213> Zea mays

 <400> 438

 gggctgcccc gtgtaggcaa gtcaaccatc atgagcgagc tttattaata tgatgcaagc 60
 tacctggttt ggtggtaatg cttatgtgat tggtgattct gcaaagcata agcagaagtg 120
 gcacgtctac tatgccacca ctgagcacc ctaggagct tgttggttact cttgagatgt 180
 gcatgactag gctggacaag aagagagctt atgttttctt caagacctct attgatgggt 240
 acacatcttg tgctaaggat atgaccaagc cttcagtgat t 281

<210> 439
 <211> 334
 <212> nucleic acid
 <213> Zea mays

 <400> 439

 gcagaagtgg cacgtctact atgccaccac tgagcacc c taaggagctt gttgttactc 60
 ttgagatgtg catgactagg ctggacaaga agagagctat gtcttcttca agacctctat 120
 tgatggttac acatcttgtg ctaaggatat gaccaagcct tcagtggcat cgtcaaggag 180
 tgcgtcgtga ccagtgggtg tagcagcaag gccatggcac tcagggtcca tgggtcgtgg 240
 gtgtggcttg ttcagcgctt actattcgcg ggagtcggcg aggtgcctgc tgaacttggt 300
 ggaagtggag tccattatga tctgacacga cctc 334

<210> 440
 <211> 349
 <212> nucleic acid
 <213> Zea mays

 <400> 440

 catctatcct ctgaagattg tcatcaagac ctgtggcact accaagctcc tgctcacaat 60
 tocaaggatc ctagagcttg tgaagagctg tctatgctct tgtgctgtga antatccccg 120
 gggacgttca tctttcttgg cgnacagcag cccccaccg gagttctccg aggagttgtg 180
 tattaaccgt actttggggg ctgaagtctg gtggcatgct tatgtgattg gagatgcagc 240
 aagaccagga cagaagtggc acatctatac gccactgagt acccagagca accatgtcac 300

cttgagatgt gcatgatggt tggacagaag aagcttcac tttttcaag

349

<210> 441
<211> 260
<212> nucleic acid
<213> Zea mays

<400> 441

tttctcttat ggngacctgg ntaagagggt gctgaggcgc ttnggtccaa ctgagcnctc 60
tgttgccgtg accatcttcg gtgatcgga caatgcgaag acctggggga cgaaactgna 120
tgctgaggcc tatgcttgca gcaacatggc tgagcaggag cttgccgatt ggtggcttgc 180
tcactatca gagcttcaact gttacggcgc aancgacctn tgnttcccag ngcaaacntc 240
gnaaattccg ggnaaantaa 260

<210> 442
<211> 447
<212> nucleic acid
<213> Zea mays

<400> 442

cgccctgca cacggcggtt ttcagactgc gatccattcc gagcttcgag gaatcgatgg 60
tcnagcttcg gtgactaaga gatcaaactc ttcaagttct acgaggctga agatcctgag 120
catctgtttg gtgaagattc ttatgccatg gaaattcatc gattgatacc gtcccttttca 180
tagtcgatca tggttcaagg agattgcatt tgtggatgct ctaatggagt cgaaaggtgg 240
taagaaagtc nnnnnnnnnn nnnnnnnnnt tcatgtacga agctcccctt ggctacaaga 300
tcgaggacgt tcgcccagcc ggaggaatca agaagttcca gactgctgct tattccaact 360
gcgtccgcca gccatcctga tatcgctca catgcaattc gcggtagagt aggattttta 420
ttcagttttc ctcttngtc ngnaggt 447

<210> 443
<211> 192
<212> nucleic acid
<213> Zea mays

<400> 443

gggctgagac ttcgannacg agccctnttg ctattncatg aatgctgnaa atnanccggg 60

gttggggana attcatntga gncnagagga cngattnagc tatgcaagct annaggtcat 120
 gggctngaac cccgactctt tntgttatgg tgacctgcct aanagggagc tgangngctn 180
 nggtccaaat ga 192

<210> 444
 <211> 376
 <212> nucleic acid
 <213> Zea mays
 <400> 444

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 tcgctgcagt taaanactnn cgtggaacgt ncatattccn tgaagcacan centtcennac 120
 acaagaactt ngntgacgag gttgccttcc tgaatngctt attcngtggn ctnaagtncn 180
 gnngcaatgc ttatgtgant ggtgactctg ccaancccg nagaagtgn cacgtctact 240
 acacnnctga gcacctgan gagcctgttn gnactctgga gatgtgcatg actgggctgg 300
 acaagancaa agcttcatgn cntcttcaag accactgnnt gatggttact cgctnatncg 360
 ccaaggatat caccaa 376

<210> 445
 <211> 502
 <212> nucleic acid
 <213> Zea mays
 <400> 445

gtgctagtgg cnangnnntn nngtatnaan gacacactcc ggtncggaat tcccgggtcg 60
 acccacgcgt ccgcgagcgc gtgggggtgc caaagaattn cctnccctac cgtcgctcgc 120
 gctcgctctgg nngggaagtt ccnggaccn ngatttngcc caattctnag gnaaaaaannc 180
 ggtttacntn ncttaaccn gggnnctctn cctntngtng ttgttaaagg ganaaaattn 240
 tnagatctgt tccgatcaa gctgcganc tcgggaatca agggttttna cacatatctt 300
 ttccatttga aaattgatnt actaaatggn gtctanaagg tggcaaaaag tctnacaata 360
 gtcgttctat natgtatgaa gctccccttn gctncagcat tnaggacgtt cgacctgcc 420
 gangcctnaa gaaatttcca gtctnctgct tactncaact tcncaaaaa accattctga 480
 tatecttttg cttnctcaat nt 502

<210> 446
 <211> 160
 <212> nucleic acid
 <213> Zea mays

<400> 446

agtnngctgca cttaaccggt nctttggcgg cctgaaatct ggtggnangg gcgggtgtga 60
 tnggagatcc agcaagacct ggncagaagt ggcacttntt nnaacgnan tnantnccna 120
 ttancaacca atggttaacc tanaaatgtg catnactgna 160

<210> 447
 <211> 487
 <212> nucleic acid
 <213> Zea mays

<400> 447

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 ccgggncggg ntcnatntnn agaannngna gtacttgtct cagcccgggg nctgctgcgt 120
 cnggtggtgg tnaaggggag aanntcgtna gatctgntcc ggatcaagcg nncganctcg 180
 ggaatcaggg gtttcacaca tanctgngnc gatcnogaat tctgatgtac taatggagtc 240
 taagggnggc aaaaagtcta gccttacnta cnctatnatg catagaagct ccccttggtc 300
 acagcattna cgaccgttcg anctgtccng aggcgtgaaa aagntccnnt acngctgctt 360
 actccaactt gcgcgaaaca anccntcctg aanantcccc ttttggtctc ctcattctaa 420
 gcactttaag gaattttaat ncttgacac ttntggantn ttnacciaan ctontctggg 480
 cctaggg 487

<210> 448
 <211> 438
 <212> nucleic acid
 <213> Zea mays

<400> 448

gagganntcg ttgagatcta agatagagaa tcgctggggc angngnctgt ggtctgctgc 60
 gtaggggtggt gntgaanggg agaagtttaa gatctgntcc acagatcacg cgtgcgcgct 120
 cgcgaancgg gggttcacac catagcttng tcggtttgaa tttgangtac taatggagtc 180

taaggggtggc aagaantcta gcagnagtcg ttccatgatg natgaagctc cccttggtcta 240
cagcattgaa gacgttcgac ctgccggaag cgccaaagaa gttncagnct tgntgcttaa 300
ctccaactgg cncaaaagaa nccatnctgn aatccccctt ggnttccccg ntctangagn 360
ttaagaattc ttttcttgca ctttnaatct taacnaatcc ccctgggctg angnttttct 420
gacaaaangaa ccaattnc 438

<210> 449
<211> 429
<212> nucleic acid
<213> Zea mays
<400> 449

ctcgattaca aactgtgtac tatgttaatg taatgacata gtttggtga tacagatcgt 60
agtgatgttg tttgaagctg tttgtatnaa ctgtntccta ctatcttgta tgaattatga 120
tgtctgatgt agtntgtatg aactatgtca tattatgttg aatgaagcta ttgcatgccc 180
ttaaaaaana aaaaccnaag annanaatga tctactccaa ntaacgagnt tatgggcttg 240
gatgccactg ntntgnntta tgggtgacct gtcaagaggg tgctnctggn ctttngccnn 300
tnagagtttt acnttgccct gaccatcttc nncggncgtg gnatgccggg acatnnggna 360
aggnacttgg tgnanaagtc tttnactggn acanctgtt cgaacangac ttntnttnaa 420
gcggnetcc 429

<210> 450
<211> 376
<212> nucleic acid
<213> Zea mays
<400> 450

ccaatatctt tactnccgcc cctgncgta attccctnct ggcncacgc atcngctga 60
agacnctn tnctacctn ccnantntg tttccactn ctttnaccac gagggngggg 120
nagattgctc ctntnctat caataangac tacaatctgg ctaatcttct ctgctnaaca 180
ngaggaatcc ccatgccatn natnaccaag ncnntagtgc ttgatnanta atacntctt 240
ctaagntcca tntncttctg aattgnntat antatctct cacantttca tantntcan 300
tnagttattc tatcaagcat ccaatccatt ctattgtata ttaaaatttn tctctctnta 360

tncatgtcaa cttcct

376

<210> 451
<211> 305
<212> nucleic acid
<213> Zea mays

<400> 451

ggaagttgga tgctatcgag gagaaggatg gagtgcttga tgagtaagac gggtttctgg 60
tgtcgatttg cttctgagtt tttttctntt ttatatcggt gcgatctcgt ggttgctcgt 120
tggttagcag ccaagccagg ctattagtag gaagaatgtc gtctataaac atgtgttcct 180
acgttgccac atgtcgagtc aatctgtata cagctagctc taggtgggtca gctgcgtcta 240
ccacaatgag catacatgta tggataaatc ttctgtgaaa gcattcgatg aataagggtt 300
gtttt 305

<210> 452
<211> 276
<212> nucleic acid
<213> Zea mays

<400> 452

atgcaagcac cttttgcgaa ggactgcaag ctggctaatac ttgtctgctt ggaggaagtt 60
ggatgctatc gaggagaagg atggagtgtc tgatgagtaa gacgggtttc tggtcgtatt 120
tgctttctgag tttttttctt ttatatcgt tgcgatctcg tggatgtcgt ttggttagca 180
gccaagccag gctattagta cgaagaatgt cgctataaac atgtgttcoo acgtgccaca 240
tgctgagtca atctgtatac agctantcta ggtggc 276

<210> 453
<211> 501
<212> nucleic acid
<213> Zea mays

<400> 453

gnaagngttt tgaatggggg ntgggngna annnnnnnct ngtaaccggtc gngngnctgn 60
cccacgnta cgcgccnttg ntagctcggc ccaaatacgt tcttngntgc tttgatggcg 120
agaatgcagc accttttgcg aaggactgca agctggctaa tcttgtctgc ttggaggaag 180

ttggatgcta tcgaggagaa ggatggagtg cttgatgagt aagacggggt tctggtgtcg 240
 atttgcttct gagttttttt ctttttatat cgttgcgac tcgtggggtg tcggtttggt 300
 tagcaagcca agccaggcta ttagtaccaa agaatgtcgn ctataaacat gtgttcctac 360
 gttgcacatc ctgagtcnaa tctgtattnc agctagctct aaggtgggtc agcttgcgtc 420
 taccacaatg agcataccat gtatggataa atcttctgtg aaagcattcg atgaataagg 480
 gttgttttca aaaaaaaaaa a 501

<210> 454
 <211> 247
 <212> nucleic acid
 <213> Zea mays

<400> 454
 gtgtcgattt gcttctgagt tntcttcttt ttatatcggt gcgatctcgt ggttgtcggt 60
 tggttagcag ccaagccagg ctattagtac gaagaatgtc gtctataaac atgtgttcct 120
 acgttgccac atgtctgagtc aatctgtata cagctagctc taggtgggtca gctgcgtcta 180
 ccacaatgag catacatgta tggtaaactt tctgtgaaag catcgntgaa taaggtngtt 240
 tcgtntt 247

<210> 455
 <211> 185
 <212> nucleic acid
 <213> Zea mays

<400> 455
 gtttggttaa gcagccaagc caggctatta gtacgaagaa tgcgtctat aaacatgtgt 60
 tctacggtg ccacatgctg agtcaatctg tatacagcta gctctaggtg gtcagctgcg 120
 tctaccacaa tgagcataca tgtatggata aatcttctgt gaaagcattc gatgaataag 180
 ggttg 185

<210> 456
 <211> 148
 <212> nucleic acid
 <213> Zea mays

<400> 456

ggcctcaaat gacgagatgt gtattggctg ctggaatg

338

<210> 460
<211> 165
<212> nucleic acid
<213> Zea mays

<400> 460

cggncttctc ggctgcagtt accatctttg gtggccgtgg gttcgccaaa tcatgggcga 60

cgggtgcaga catcgattcc tacatgtgcg gtgatcctgt agagcaagag cttcctgggtg 120

gcggtctgct gatgtaccag agctttactg ctgttccctc tggct 165

<210> 461
<211> 293
<212> nucleic acid
<213> Zea mays

<400> 461

agcgcgccta cctattcctt ctcccagccg tcgagctgcc caaccagag caccagcagc 60

cctcaccttc ttctctcgg cgggtccttc aaacgagccc gctcgtcttg gagccgcgcg 120

cggctctggt ctggacgaac gttcaaggag attgcatttg tggatgctct aatggagtcg 180

aaaggtggca agaagtcnnn nnnnnnnnnn nnnnnnttca tgtacgaagc tccccttggc 240

tacaagatcg aggacgttcg cccagccgga ngaatcaaga agttccagac tgc 293

<210> 462
<211> 289
<212> nucleic acid
<213> Zea mays

<400> 462

gacctattcc ttctcccagc cgtcgagctg cccaaccag agcaccagca gccctacacc 60

ttcttctctt acggccggtc cttcaaacga gcccgctcgt acttggagcc gccgccggta 120

ctcgttctgg acgaacgttc aaggagattg catttgtgga tgcctaatg gactcgaaag 180

gtggcaagaa gtcnnnnnnn nnnnnnnnnn nnttcatgta cgaagctccc cttggctaca 240

agatcgagga cgttcgccca gccggagaat caagcagttc cgactgatg 289

<210> 463

<211> 264
 <212> nucleic acid
 <213> Zea mays
 <400> 463
 attcctttctc ccagccgtcg agctgcccac cccagagcac cagcagccct cacctttcttc 60
 ctctcggccg gtccttcaaa cgagcccgtc cgtcttggag ccgccgccgg tctcgttctg 120
 gacgaacgtt caaggagatt gcatttgtgg atgctctaata ggagtcgaaa ggtggcaaga 180
 agtcnnnnnnn nnnnnnnnnn nnttcatgt acgaagctcc ccttggctac aagatcgagg 240
 acgttcgccc agccggagga atca 264

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<210> 464
 <211> 571
 <212> nucleic acid
 <213> Zea mays
 <400> 464
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 cacgcgtccg gtccggtggc aatgcttatg tgattggtga ctctgcgaag cccgggcaga 120
 agtggcacgt ntactacgcc gctgagcacc ctgaggagcc tgtcgttact ctggagatgt 180
 gcatgactgg gctggacaag aagaaagctt ctgtcttctt caagaccact gctgatgggt 240
 actcgtctgt cgccaaggag atgaccaagc tctctggtat ctccgacatc atcccagaga 300
 tggagatctg tgacttcgac ttcgagccct gtggctactc catgaatgct gttcatggcc 360
 ctgctttgtc gaccattcat gtgaccccag aggacggctt cagctatgca agctacgagg 420
 tcatgggctt naaccggggc tctttctctt atggngacct ggtaaaaaag ggtctnaagt 480
 nctttggtca actgaattct ntggtgccgt gaccatntn gngaannnga caaatgcnaa 540
 aaccttgggg accaaactgg attcttaagg c 571

<210> 465
 <211> 522
 <212> nucleic acid
 <213> Zea mays
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 agagggctcg nngtngana ggngnnnnag gnantttttg naattccgaa ttcccgggtn 60

gaccacgcg tccgctnatg tcttatctga gtcgagcctg tttgtctacc cgtacaagat 120
 tgtgatcaag nggtgtggaa gtacaaagct tctgctcgct attccgagga tccttgaact 180
 tgctgaagag ctctgtttgc cactcgctgc agttaaatat tcccgtggaa cgttcatatt 240
 ccctgaagca caaccttccc cacacaagaa cttcgctgac gaggttgccct tcctgaatcg 300
 cttcttcggt ggctcaagt ccggtggcaa tgcttatgtg attggtgact ctgcgaagcc 360
 cgggcagaag tggcacgtct actacgccgc tgagcaccct gaggagcctg ttgttactct 420
 agagatgtgc atgactgggc tggacaagaa gaaagcttct gtcttcttca agaccactgc 480
 tgatggttac tcgctgtgcg ccaaggagat gaccaactct ct 522

<210> 466
 <211> 515
 <212> nucleic acid
 <213> Zea mays

<400> 466
 gagnttgntt cnacgnaggg gggatntaac ggaaangctt gtaccggtcc ggaattcccg 60
 ggtcgacca cgcgtccgat tccctgaagc acagccttcc ccacacaaga acttcgctga 120
 cgaggttgcc tncctgaatc gcttcttcgg tggcctcaag tccggtggca atgcttatgt 180
 gattggtgac tctgcgaagc ccgggcagaa gtggcacgtc tactacgccg ctgagcacc 240
 tgaggagcct gttgttactc tagagatgtg catgactggg ctggacaaga agaaagcttc 300
 tgtcttcttc aagaccactg ctgatggta ctcgctgtgc gccaaaggaga tgaccaagct 360
 ctctggatc tcggacatca tcccagagat ggagatctgt gacttcgact tcgagccctg 420
 tggctactcc atgaatgctg gtcatggcct gcttggcgac cattcatgtg accccaanag 480
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<210> 467
 <211> 531
 <212> nucleic acid
 <213> Zea mays

<400> 467
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cctgcaagtt gctgctggtg cccctcccc tgtctctgtg attgggttcg agggatttga 180
gaagcgctt gagatcagct tctctgaggc acctgtcttg gctgaccca gcggaagggg 240
actgctgctg ctctcgctg cccagatcga ctctgttctt gaccttgctc ggtgcaccat 300
tgtgtccgag ctctcaaagc aggacttcga ctcttatgtc ttatctgagt cgagcctgtt 360
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tccgaggatc cttgaacttg ctgaagagct cctgttgcca ctgctgcaa gttaaatact 480
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<210> 468
<211> 463
<212> nucleic acid
<213> Zea mays
<400> 468

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ttgagaagcg ccttgagatc agcttctctg aggcacctgt cttggctgac cccagcggaa 180
ggggactgctg tgcgctctcg cgtgcccaga tcgactctgt tcttgacctt gctcggtgca 240
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<210> 469
<211> 474
<212> nucleic acid
<213> Zea mays
<400> 469

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accaagctct ctggtatctc ggacatcatc ccagagatgg agatctgtga cttcgactcg 180
agccctgtgg ctactccatg aatgctgttc atggccctgc tttgtcgacc attcattgac 240

cccagaggac ggcttcagct atgcaagcta cgaggatcatg ggctttaacc cgggcctttc 300
tcttatgggtg acctgggttaa gaggggtgctg aggtgctttg gtccaactga gtttctgttg 360
ccgtgaccat cttcggtgat cgggacaatg cgaagacctg ggggacgaaa ctgatgctga 420
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<210> 470
<211> 304
<212> nucleic acid
<213> Zea mays
<400> 470

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tctctgggtat ctgggacatc atcccagaga tggagatctg tgacttcgac ttcgagccct 120
gtggtactc catgaatgct gttcatggcc ctgctttgtc gaccattcat gtgaccccag 180
aggacggctt cagctatgca agctacgagg tcatgggctt taaccggggc tctttctctt 240
atggtgacct gggttaagagg gtgctgaggt gctttgttcc aactgagttc tctgttgccg 300
tgac 304

<210> 471
<211> 450
<212> nucleic acid
<213> Zea mays
<400> 471

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cttcaaccgg ggctctttct cttatgggtga cctgggttaag aggggtgctga ggtgctttgg 180
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aagagtgatg ctccctggga agctgatctg 450

<210> 472

<400> 472

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<400> 473

<400> 474

165

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agcggaaggg gactgcgtgc gctctcgcgt gccagatcg actctgttct tgaccttgc 240
cggcgcacca ttgtgtccga gctctcaaac gaggacttcg actcttatgt cttatctgag 300
tcgagcctgt ttgtctaccc atacaagatt gtgatcaaga cctgtggaac tacaaaagctt 360
ctgctcgcta ttccgaggat cttgaacttg ctgaaaaact cctggtgcca ctgctgcaa 420
gttaaatact nccgtggaac gttcat 446

<210> 475
<211> 294
<212> nucleic acid
<213> Zea mays

<400> 475
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tctgctcgct attccgagga tcttgaact tgctgaagag ctctgttgc cactcgctgc 180
agttaaatac tcccgaggaa cgttcatatt cctgaagca cagccttccc cacacaagaa 240
cttgcgtgac gaggttgctt tctgaatcg cttcttcggt ggcctcaagt ccgg 294

<210> 476
<211> 314
<212> nucleic acid
<213> Zea mays

<400> 476
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ntgngatgg ttannnnngn gtgcgcnaag gngatgacca agctctctgg tatctcggac 120
atcatcccag agatggagat ctgtgacttc gacttcgagc cctgtggcta ctccatgaat 180
gtgtttcatg gccctgcttt gtcgaccatt catgtgacct cagaggacgg cttcagctat 240
gcaagctacg aggtcatggg ctttaacccg ggctctttct cttatggtga ctgggtaaga 300
gggtgctgag gtgt 314

<210> 477

<211> 309
 <212> nucleic acid
 <213> Zea mays

 <400> 477

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 ctttgactct tatgtcttat ctgagtcgag cctgtttgtc taccctaca agatttgtgat 180
 caagacctgt ggaactacaa agcttctgct cgctattccg agnatccttg aacttgctga 240
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 agcacaacc 309

<210> 478
 <211> 245
 <212> nucleic acid
 <213> Zea mays

 <400> 478

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 tgtgcgcca ggagatgacc aagctctctg gtatctcgga catcatccca gagatggaga 120
 tctgtgactt cgacttcgag ccctgtggct actccatgaa tgctgttcat ggccctgctt 180
 tgtogaccat tcatgtgacc ccagaggacg gcttcagcta tgcaagctac gaggtcatgg 240
 gcttt 245

<210> 479
 <211> 592
 <212> nucleic acid
 <213> Zea mays

 <400> 479

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 cgggtcgacc cagcgcgccg gtcagggcnc aatgctnatg tgattgggga ntctgcnaag 120
 cccggtcaga agtggcacgt gtactaggcc gctnagcacc ntgaggagcc tgctgttact 180
 ctggagatgt gcatgactgg gctggacaag aagaaagctt ctgtattgtt caagaccact 240
 gctgatgggt actngctgtg cgccaaggag atgaccaagc tctctggtat ntnggacatc 300

attccagaga tggagatctg tgacttcnac ttngagccct gtgggctact nnatgaatgc 360
 tgttcatggc cctgnttttg tcgaccattc atgtgacccc aaaaggacgg nttttaactn 420
 ntgcaagcta cgaggttatg ggctttaaac ccnggncttt tttnnnatgg gggacctnng 480
 taaaaagggg tgctgaagtt ncttttggtc caactgaatt nntntggtgn ctgagacnat 540
 ttnnngtgat cgggacnant tcgantaact tnggggnchna aaacttagtg nt 592

<210> 480
 <211> 260
 <212> nucleic acid
 <213> Zea mays

<400> 480

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 tccggtggca atgcttatgt gattggtgac tctgcgaacc cgggcagaag tggcacgtct 180
 actacgccgc tgagcaccct gaggagcctg ttgttactct agagatgtgc atgactgggc 240
 tggacaagaa gaaagcttct 260

<210> 481
 <211> 249
 <212> nucleic acid
 <213> Zea mays

<400> 481

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 ccgggcagaa gtggcacgtc tactacgccg ctgagcacc ctgaggagcct gttgttactc 180
 tagagatgtg catgactggg ctggacaaga agaaagcttc tgtcttcttc aagaccactg 240
 ctgatggtt 249

<210> 482
 <211> 258
 <212> nucleic acid
 <213> Zea mays

<400> 482

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gctttgtcga ccattcatgt gacccagag gacggcttca gctatgcaag ctacgaggtc 120
atgggcttta acccgggctc tttctcttat ggtgacctgg ttaagagggg gctgaggtgc 180
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acctggggga cgaaactg 258

<210> 483
<211> 198
<212> nucleic acid
<213> Zea mays

<400> 483

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gatctgtgac ttcgacttcg agccctgtgg ctactccatg aatgctgttc atggccctgc 120
tttgtcgacc attcatgtga cccagagga cggttcagc tatgcaagct acgaggtcat 180
gggcttcaac ccgggctc 198

<210> 484
<211> 246
<212> nucleic acid
<213> Zea mays

<400> 484

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cgggctcttt ctcttatggg gacctgggta agaggggtgt gaggtgcttt ggtccaaactg 120
agttctctgt tgccgtgacc atcttcgggtg atcgggacaa tgcgaagacc tgggggacga 180
aactggatgc tgaggcctat gcttgcagca acatgggtga gcaggtgctg ccgtttggng 240
gcttgc 246

<210> 485
<211> 302
<212> nucleic acid
<213> Zea mays

<400> 485

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gaggatcctt gaacttgctg aagagctcct gtngccaacna cgctgcagtn aaatacnccc 120
 cgtggaacgt tcatattccc tgaagcacag ccttccccac acaagaactt cgctgacgag 180
 gttgccttcc tgaatcgctt cttcgngngc ctcaagtccg gnggcaatgc ttatgtgatt 240
 ggngacnctg cgaacccggg cagaagtngc agtctanacg ccgctgagca cnctgaggag 300
 cc 302

<210> 486
 <211> 524
 <212> nucleic acid
 <213> Zea mays

<400> 486

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 gaanccctga aagcacagcc ttccccacac aagaacttcg ctgacgaggt tgccttcctg 120
 aatccgcttc ttcggtggcc tcaagtccg tggcantgct tatgtgattg gtgactctgc 180
 gaagcccggg cagaagtggc acgtctaacta cgccgctgaa gcaccctgag gagcctgttg 240
 ttactctaag agatgtgcat gactgggctg gacaagaaga aagcttctgt cttcttcaag 300
 accaactgct gatggttact cgctggtgcg ccaaggagat gaccaaagct ctctggtatc 360
 tcgggacatc atcccaagag atggagatct gtgactcgac ttcgaagccc tgtggctact 420
 ccatgaatgc tgttcaatgg ccctgctttt gtgaancat tcatgtganc ccaagaggga 480
 cggctttcag ctattccaag cttacnaagg tcatgggctt taac 524

<210> 487
 <211> 178
 <212> nucleic acid
 <213> Zea mays

<400> 487

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 ccagaggang gcttcagcta tgcaagcnac gaggtcatgg gctttaacnn gggtcttt 178

<210> 488
 <211> 238
 <212> nucleic acid

<213> Zea mays

<400> 488

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cctgggtaag aggggtgctga ggtgctttgg tccaactgag ttctctgttg ccgtgaccat 120
cttcggtgat cgggacaatg cgaagacctg ggggacgaaa ctggatgctg aggcctatgc 180
ttgcagcaaa atgggttaaac agttctgccg ttgggggnntt cccacannaa aanaactt 238

<210> 489

<211> 304

<212> nucleic acid

<213> Zea mays

<400> 489

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gacaaaaaga tatcaaagac caattcccgt cttectcccg tggctgtctg tcattctggct 120
ctgaaacgat ggcagtcctg caagttgctg ctgctgcccc tccccctgtc tctgtgattg 180
ggttcgaggg atttgagaag cgccttgaga tcagcttctc tgaggcacct gtcttggtg 240
accccagcgg aaggggactg cgtgcgctct cgcgtgccca gatcgactct gttcttgacc 300
ttgc 304

<210> 490

<211> 229

<212> nucleic acid

<213> Zea mays

<400> 490

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tgctgctgct gcccctcccc ctgtctctgt gattgggttc gagggatttg agaagcgct 120
tgagatcagc ttctctgagg cacctgtctt ggctgacccc agcgggaagg gactgcgtgc 180
gctctcgcgt gccagatcg actctgttct tnaactgctc ggtgcacca 229

<210> 491

<211> 290

<212> nucleic acid

<213> Zea mays

<400> 491

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ttgagcaggt gctgccgttt ggtggcttgc tcatctatca gagcttcact gttacggccg 180
aaacgaccca tgggtcgccg aggtcagtec tgcattgactt cgctgggtgac attgtcaaga 240
atcgctcaga gaggatgct cccgtgggaag ctgatgctgt ggatgcggng 290

<210> 492

<211> 289

<212> nucleic acid

<213> Zea mays

<400> 492

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gagctcctgt tgccactcgc tgcagttaaa tactcccgtg gaacgttcat attccctgaa 180
gcacaacctt cccaacaca agaacttcgc tgacgagggt tgcctnctg aatcgctct 240
cgggtgggctc aangtcgggtg ggaaagctan tgtgattgtn annnngcga 289

<210> 493

<211> 318

<212> nucleic acid

<213> Zea mays

<400> 493

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tgagcaggtg ctgccgtttg gtggcttgct catctatcag agcttcactg ttacggccga 180
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tgtaaagaag atgnaatg 318

<210> 494

<211> 310

<212> nucleic acid

<213> Zea mays
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 tcccgtggct gtctgtcatc tggctctgaa acgatggcag tcctgcaa at tgctgtgct 180
 gccctcccc ctgtctctgt gattgggttc gagggatttg agaagcgcct tgagatcagc 240
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 ccagatoga 310

<210> 495
 <211> 137
 <212> nucleic acid
 <213> Zea mays
 <400> 495
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 acctggggga cgaaact 137

<210> 496
 <211> 111
 <212> nucleic acid
 <213> Zea mays
 <400> 496
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<210> 497
 <211> 302
 <212> nucleic acid
 <213> Zea mays
 <400> 497
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atggagtcca aggggtggcaa gaagaagtct agcagtagtc gttcctccct gatgtacgaa 180
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ag 302

<210> 498
<211> 501
<212> nucleic acid
<213> Zea mays
<400> 498

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tcnaactgcn cgaggaagcc atctgatata ctctgtctc atccccatcc taatagccgt 420
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<210> 499
<211> 284
<212> nucleic acid
<213> Zea mays
<400> 499

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gacaaaaaga tatcagagac cacctcccgt ggctgtctgc catctggctc tgaaacgatg 180
gcagtcctac aagctgtctc cctccccct gtctctgtga ttggattcga gggatttgag 240
aagcgccttg agatcagctt ctctgaggca cctgtcttgg ctga 284

<210> 500

<211> 166
 <212> nucleic acid
 <213> Zea mays

<400> 500

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<210> 501
 <211> 440
 <212> nucleic acid
 <213> Zea mays

<400> 501

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 aagaagaagt ctagcagtag tcgttctctc ctgatgtacg aagctccctt tggctacagc 180
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<210> 502
 <211> 300
 <212> nucleic acid
 <213> Zea mays

<400> 502

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 gaggaagcca tctgatata tctgtcgtca tccccatcct agtagcgtag aaacctcttg 180
 cattttcatt ttgatctccc taatctctcc ggctagctgc tttccagtga ccaaaagata 240
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<210> 503
 <211> 283
 <212> nucleic acid
 <213> Zea mays

 <400> 503

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 ctggctctga aacgatggca gtcttacaag ctgctgcccc tccccctgtc tctgtgattg 180
 gattcgaggg atttgagaag cgccttgaga tcagcttctc tgaggcacct gtcttggtg 240
 accccagcgg aaggggactg cgtgcgctct cgcgtgccca gat 283

<210> 504
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 <212> nucleic acid
 <213> Zea mays

 <400> 504

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 cgctgggttc tgcgagangt cggggccttc gtgggatctt gaaaattgaa tgtttctaag 180
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 <400> 505

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 gcagcagaat tcgtcgaggg ccgctgggtg ctgcgagang tccgggcctt cgtgggatct 180
 tgaaaattga atgtttctaag ggagtccaag ggtggcaaga agaagtctag cagtagtcgt 240
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<211> 283
<212> nucleic acid
<213> Zea mays

<400> 506

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cgctgggttgc tgcgagangt ccgggccttc gtgggatctt gaaaattgaa tgttctaata 180

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ccccttggct acagcatcga ggatatccgc cctgcaggcg gca 283

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<400> 507

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cagcagaatt cgtcgagggc cgctgggttgc tgcgagaggt ccgggccttc gtgggatctt 180

gaaaattgaa tgttctaata gagtccaagg gtggcaagaa gaagtctagc agtagtcggt 240

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gcat 304

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<212> nucleic acid
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<400> 508

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cgagcagcag aatttgctga gggccgctgg ttgctgcgag aggtccgggc cttcgtggga 180

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 cgttcctccc tgatgtacga agctcccctt ggctacagca tcgaggatat ccgcctgcag 300
 gcggcatcaa gaagttctcc gctgcttact cgaactgcgc gaggaaacca tntgatatac 360
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<400> 509

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 aangaaagaa atctagcagt agtcgttctt tctgatgta cgaaacttcc nttggetnca 300
 acatcgagga tattcgccct tgaagcngca ttaaanantt nttcnntngt tactnnaant 360
 ttncangaa accattctta aatnctttgt ngnattccca tcctaatacg tttnaaacan 420
 ctgaatttca attgacttct naaaanctag ggt 453

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<400> 510

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 ccgctgggtg ctgcgagang tccgggcctt cgtgggatct tgaaaattga atgttctaata 180
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<400> 512

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 agatctcttg actcacgaca cgagcagcag aattcgctga gggccgctgg ttgctgcgag 180
 aggtccgggc ctctgtggga tcttgaaaat gaatgttcta atggagtcca agggtgggca 240
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 cgaggat 307

<210> 513
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 <212> nucleic acid
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 cgagcagcag aattcgctga gggccgctgg ttgctgcgag aggtccgggc ctctgtggga 180
 tcttgaaaat tgaatgttct aatggagtcc aagggtggca agaagaagtc tagcagtagt 240
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 gcggcat 307

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 <212> nucleic acid
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 gagcagcaga attcgtcgag ggccgntggg tgctgcgaga ggtccggggc ttcgtgggat 180
 cttgaaaatt gaatgttcta atggagtcca aggggtggca gaagaagtct agcagtagtc 240
 gttcctccct gatgtacgaa gtccctggc taca 274

<210> 515
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 <212> nucleic acid
 <213> Zea mays

 <400> 515

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 agcagaattc gtcgaggggc gctgggttgc gcgagaggtc cgggccttcg tgggatcttg 180
 aaaattgaat gttctaattg agtccaaggg tggcaagaag aagtctagca gtagtcgttc 240
 ctccctgatg tacgaagctc cccttggtc cagcatcgag gatatccgcc 290

<210> 516
 <211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 516

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 gcgggttaggg ttgggagatc tcttgactca cgacacgagc agcagaattc gtcgaggggc 180
 gctgggttgc gcgagaggtc cgggccttcg tgggatcttg aaaattgaat gttctaattg 240
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<210> 517
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 tctcttgact cagcacacga gcagcagaat tcgtcgagggt ccgctgggtg ctgcgagagg 180
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 aggat 305

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 aattgaatgt tctaattggag tccaagggtg gcaagaagaa gtctagcagt agtcgttcc 240
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<210> 520

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<210> 521

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agatctcttg actcacgaca cgagcagcag aattcgtcga gggccgctgg ttgctgcgag 180

agggtccgggc cttcgtggga tcttgaaaat tgaatgttct aatggagtcc aaggggtggca 240
agaagaagtc tagcagtagt cgttcctccc tgatgtacga agctccccctt ggctacagca 300
tcgaggatat ccg 313

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<400> 523

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agatctcttg actcacgaca cgagcagcag aattcgtcga gggccgctgg ttgctgcgag 180
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agaagaagtc tagcagtagt cgttc 265

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<213> Zea mays

<400> 524

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gagatcncctt gactcacgac acgagcagca gaattcncg agggccgctg gttgcngcga 180
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aagaagaagt ctagcagtag tcgttcctcc ct 272

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<212> nucleic acid
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<400> 525

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gcttactcca actgcgcgaa gaagccatcc tgatatccct tttggcttcc tcattctagt 180
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 ttgtggaccc tcatggggcg tggtttgctg gccctctnca gggcccanat tgactctgtt 480
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<210> 526
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 <212> nucleic acid
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<400> 538

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<213> Zea mays

<400> 539

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 <213> Zea mays
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<400> 545

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<400> 548

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<211> 512

<212> nucleic acid

<213> Zea mays

<400> 550

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actaccaagc tcctgctcaa tantcnaaga at 512

<210> 551

<211> 448

<212> nucleic acid

<213> Zea mays

<400> 551

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gcatgactgg tctggacaaa gaaganagct tctgtctttt tcaaagacta atgctgatgg 180
gaacacaaca tgtgccaagg aatgacaaa gctctctggc atctctgaaa tcatccccga 240
gatggagatc tgcgattttg actttgaacc ctgtggctac tccatgaatg cgatccatgg 300
ctctgcattc tccacaatcc atgtgacgcc cgaggacggt ttcagctatg ccaagttatg 360
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<210> 552
<211> 384
<212> nucleic acid
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<400> 552
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ggacgttcat ctttctgtgt gcacagccag cccccacag gagcttctct gaggaagttg 180
ctgcacttaa ccgctacttt ggcggcctga aatctggtgg taatgcttat gtgattggag 240
atccagcaag acctggacag aagtggcacg tcttctacgc cactgagtac ccagagcaac 300
caatggntaa ccttgagatg tgcattgact gtctggacaa gaaagaaagc ttctgtcttt 360
ttcaaagact aaatgctgat ggga 384

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<211> 469
<212> nucleic acid
<213> Zea mays

<400> 553
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ccacngatcc atgtgacgcc tgaggatggg ttcagctacg ccagttatga ggttatgggc 180
ttggatgcca ctgccctgtc ttacggtgac cttgtcaaga gggccttgg gtgcttcggc 240
ccctcagaat tttctgtcgc cgtgaccatc ttcggcgggc ggngccaagc tgggacatgg 300

ggaaaggaac ttggtgcgga ggcttatgac tgcaacaaca tggtcgagca ggagctgcct 360
ggaggtggga tcctcatcta ccaaagcttc ttgtgctgnt gaaanacccc gttgctagct 420
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<212> nucleic acid
<213> Zea mays
<400> 554

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tgctttggcc cctcagagtt ttccgttgcc gtgaccatct tcggcgggcg tggccatgcc 180
gggacatggg gaaaggcact tgggtgcagag gtctatgact gcaacaacat ggtggagcag 240
gagctgcctg gaggcgggct cctcgtgtac cagagcttct gtgctgctga agacgctgtt 300
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<210> 555
<211> 435
<212> nucleic acid
<213> Zea mays
<400> 555

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tggagatctg cgattntgac ttccnaccct gtggctactc catgaatgag atccatggct 180
ctgcattctc cacaatccat gtgacgcccg agnccggttt cagctacncc agttacgagg 240
ttatgggctt ggatgccact gntntgtctt atggcgacct tgtcaagagg gtccttcggt 300
gctttgcccc tcagagtttt acgangccgt gaccatcttc ngcngaagtg nccatgcctg 360
gacatggggg aaaggcactt ggtgccnagg cctattnctg nnacancatg gtggaacang 420
agcngccttg angca 435

<210> 556
<211> 334
<212> nucleic acid

<213> Zea mays

<400> 556

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gaacgtggag agtgctcctc ctctatgaa gaaggactac aaactagcta atcttctctg 180
ctgggaggag gaagcggatg ccatggagga gaaggcggga gtgcttgatg agtaagacgg 240
gttctgtgtg tcgatttgct tctgagttgt ttattttata tcgtcgcaat ttcgtgggtg 300
tcgtttgggtt attctgtgaa gcagccaagc cagg 334

<210> 557

<211> 404

<212> nucleic acid

<213> Zea mays

<400> 557

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gggtcgaccc acgcgtccgc tcatatgtcc tttctgagtc aagcttgttt atctatcctc 120
tgaagattgt catcaagacc tgtggcacta ccaagctcct gctcaccatt ccaagaatcc 180
ttgagcttgc tgaagagctg tctatgccac ttgctgctgt gaagtactcc cgtgggacgt 240
tcattcttcc tggcgcacag ccagccccc acaggagctt ctctgaggaa gttgctgcac 300
ttaaccgcta ctttggcggc ctgaaatctg gtggaatgc ttatgtgatt ggagatccag 360
caagacctgg acagaagtgg cacgtcttct acgccactga gtac 404

<210> 558

<211> 430

<212> nucleic acid

<213> Zea mays

<400> 558

atcgggtttg agggctatga gaagcgctt gagatcacat tctctgaggc acctgtcttt 60
gtggaccctc atgggcgtgg tttgcgtgcc ctctccaggg ccagattga ctctgttctg 120
gatcttgcac ggtgcacaat tgtgtccgag ctctncaaca aggatttcga ctcatatgtc 180
ctttctgagt caagcttggtt tatctatcct ctgaagattg tcatcaagac ctgtggcact 240

accaagctcc tgctcaccat tccaagaatc cttgagcttg ctgaagaact gtctatgcca 300
 cttgtctgntg tgaaagtact cccgtgggga cgttcatctt tcctggcgca caagncaggn 360
 cccacaagg agcttntttt gaaggaagtt gcttgnactt taaccggta atttggggg 420
 ccttgaaant 430

<210> 559
 <211> 319
 <212> nucleic acid
 <213> Zea mays
 <400> 559

gtgcatgact ggtctggaca agaagaaagc ttgtgtcttt ttcaagacta atgctgatgg 60
 gaacacaaca tgtgccaagg aaatgacaaa gctctctggc atctctgaaa tcatccccga 120
 gatggagatc tgcgattttg acttogaacc ctgtggctac tccatgaatg cgatccatgg 180
 ctctgcattc tccacaatcc atgtgacgcc cgaggacggg ttcaactacg ccagttacga 240
 ggttatgggc ttggatgcca ctgctctgtc ttatgggtgac cttgtcaaga gggctcctcg 300
 gtgctttggc ccctcagag 319

<210> 560
 <211> 346
 <212> nucleic acid
 <213> Zea mays
 <400> 560

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 gcaaccaatg gttaaccttg agatgtgcat gactggctctg gacaagaaga aagcttgtgt 120
 ctttttcaag actaatgctg atgggaacac aacatgtgcc aaggaaatga caaagctctc 180
 tggcatctct gaaatcatcn ttnagatgga gatctgcat tttgacttcg aaccctgtgg 240
 ctactccatg aatgogatcc atggctctgc attctccaca atccatgtga cgcccaggga 300
 cggtttcagt acgccattac gaggttatgg gcttggatgc catgct 346

<210> 561
 <211> 317
 <212> nucleic acid
 <213> Zea mays

<400> 561

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aaagctctct ggcacatctg aatcatccc cgagatggag atctgogatt ttgacttcga 180
accctgtggc tactccatga atgcgatcca tggctctgca ttctccacaa tccatgtgac 240
gcccaggagc ggtttcagct acgccagtta cgaggttatg ggcttgatg ccactgctct 300
gtcttatggg gaccttg 317

<210> 562

<211> 342

<212> nucleic acid

<213> Zea mays

<400> 562

cagaggaggt ttcagctatg ccagttacga ggttatgggc ttggatgcc ctgctctatc 60
ttatggtgac cttgtcaaga gggtcctccg gtgctttggc ccctcggagt tttccgttgc 120
cgtgaccatc ttcggcgggc gtggccatgc cgggacatgg ggaaaggcac taggtgcaga 180
ggcttatgac tgcaacaaca tgggtggagca ggagctgcct ggaggcgggc tctcgtata 240
ccagagcttc tgcgctgctg aagacgctgt tgctacctg cccaaatctg ttttccactg 300
ctttgacggc gagaacgtgg agagtgtcct cctcctatga ag 342

<210> 563

<211> 314

<212> nucleic acid

<213> Zea mays

<400> 563

gctgcttact ccaactgcgc gaagaagcca tctgatatc ccttttggct tctcattct 60
agtagtttag gattttcttt ctgacacttt gattctgacc aatctctctg gctgctgct 120
tctgataat cgaccagttc ccagtccttg ctcttgacac tctccctcc atctccagca 180
ttgtgttctg attcaactgc tccaatggct gttctttctg ctgctgatgc tccccggtc 240
tcagctatcg ggtttgaggc ctatgagaag cgccttgaga tcacattctc tgaggcacct 300
gtctttgtgg accc 314

<400> 566

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tccagtctgc tgcttactcc aactgcgcga agaagccatc ctgatatccc ttttggttc 120
cncattctag tagtttagga tttcttttct gacactttga ttctgaccaa tctctctggc 180
ctgctnttcc tgataatcga ccagttcccc agtcttgctc cttgcaactcc tccctcctcc 240
atctccagea ttgtgttctg attcacctgc tccaatggct gttctttctg ctgctgatgc 300
ttccccggtc tcagctatcg ggttgagggc tatgagaagc gccttg 346

<210> 567

<211> 303

<212> nucleic acid

<213> Zea mays

<400> 567

ccaagctcct gctcaccatt ccaagaatcc ttgagcttgc tgaagagctg tctatgccac 60
ttgctgctgt gaagtactcc cgtgggacgt tcattcttcc tggcgcacag ccagccccc 120
acaggagctt ctctgaggaa gttgctgcac ttaaccgcta ctttggcggc ctgaaatctg 180
gtggtaatgc ttatgtgatt ggagatccag caagacctgg acagaagtgg cacgtcttct 240
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aca 303

<210> 568

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 568

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tgaccttgtc aagaggggtcc tccggtgctt tggccctcgc gagttttccg ttgcctgac 180
catcttcggc gggcgtggcc atgccgggac atggggaaag gcactagggtg cagaggtcta 240
tgactgcaac aacatggtgg agcaggagct gcctggaggc gggctcctcg tataccagag 300
cttctgcgct gctgaa 316

<210> 569
 <211> 313
 <212> nucleic acid
 <213> Zea mays

 <400> 569

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 ctgctcacca ttccaagaat ccttgagctt gctgaagagc tgtctatgcc acttgctgct 180
 gtgaagtact cccgtgggac gttcatcttt cctggtgcac agccagcccc ccacaggagc 240
 ttctctgagg aagttgctgc acttaaccgc tactttggcg gcctgaaatc tggtggtaat 300
 gcttatgtga ttg 313

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<210> 570
 <211> 512
 <212> nucleic acid
 <213> Zea mays

 <400> 570

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 gaccacggcg tccgcggaag cgtggggcgat tcttaccaat ccccttggtc tgctgctttc 120
 ctgataatcg accagttccc cagtcttggc cctgcagtc ctccctcctc catctccagc 180
 gttgtgttct gactcacctg caccaatggc tgttctttct gctgctgggtg ctcccccggc 240
 ctgagctatc gggtttgagg gctatgagaa gcgccttgag atcacattct ctgaggcacc 300
 tgtctttgtg gacccccatg gcagcgggtt gcgtgcctc tccaggtccc agattgactc 360
 tgttctggat cttgcacggt gcacgatcgt gtccgagctc tccaacaagg attttgactc 420
 ctatgttctc tccgagtcaa gcttgtcatc tatcctctga agattgncat caagacctgt 480
 ggcactacca acttctggta caanttccag ga 512

<210> 571
 <211> 314
 <212> nucleic acid
 <213> Zea mays

 <400> 571

tnggagtcta aggggtggcaa aaagtctagc agtagtcggt ccatgatgta tgaagctccc 60
 cttggctaca gcattgagga cgttcgacct gccggagngt gaagaagttc cagtctgctg 120
 cttactccaa ctgcgcgaag aagccatcct gatataccctt ttggcttccct cattctagta 180
 gtttaggatt tcttttctga cactttgatt ctgaccaatc tctctggcct gctgcttccct 240
 gataatcgac cagttcccca gtcttgctcc ttgcaactcct cctcctcca tctccagcat 300
 tgtgttctga ttca 314

<210> 572
 <211> 422
 <212> nucleic acid
 <213> Zea mays

<400> 572

cgccagttat gaggttatgg gcttgatgc cactgccctg tottacgggtg accttgtcaa 60
 gagggctcctt ggggtgcttcg gccctcggga attttctgtc gccgtgacca tcttcggcgg 120
 gcggngccaa gctgggacat ggggaaagga acttggtgcg gaggttatg actgcaacaa 180
 catggtcgag caggagctgc ctggaggtgg gatcctcatc taccagagct tctgtgctgc 240
 tgaagaogcc gttgctagct cgcccaaatc cgttcttcgc tgctttgatg gcgagaatgc 300
 agcacotttt gcgaaggact gcaagctggc taatcttgtc tgcttgaggg aagttggatg 360
 ctatogagga gaaagatgga gtgcttgatg agtaaaacng gtttcttggt gncgatttgc 420
 tt 422

<210> 573
 <211> 295
 <212> nucleic acid
 <213> Zea mays

<400> 573

tgaatcgat ccatggctct gcattctcca caatccatgt gacgcccag gacggtttca 60
 gctacgccag ttacgaggtt atgggcttgg atgccactgc tctgtcttat ggtgaccttg 120
 tcaagagggt cctccggtgc tttggccctt cagagttttc cgttgccgtg accatcttcg 180
 gcgggctggt ccatgccggg acatggggaa aggcacttgg tgcagaggtc tatgactgca 240
 acaacatggt ggagcaggag ctgcctggag gcgggctcct cgtgtaccag agctt 295

<210> 574
 <211> 295
 <212> nucleic acid
 <213> Zea mays

 <400> 574

 ggcggtggttt gcgtgccctc tccagggccc agattgaactc tgttctggat cttgcacggt 60
 gcacaattgt gtctgagctc tccaacaagg atttcgactc atatgtcctt tctgagtcaa 120
 gcttgtttat ctatcctctg aagattgtca tcaagacotg tggcactacc aagctcctgc 180
 tcaccattcc aagaatcctt gagcttgctg aagagctgtc tatgccactt gctgctgtga 240
 agtactcccg tgggacgttc atctttcctg gcgcacagcc agccccccac aggag 295

<210> 575
 <211> 299
 <212> nucleic acid
 <213> Zea mays

 <400> 575

 gtcctctctg agtcaagctt gtttatctat cctctgaaga ttgtcatcaa gacctgtggc 60
 actaccaagc tctgtctcac cattccaaga atccttgagc ttgctgaaga gctgtctatg 120
 ccacttgctg ctgtgaagta ctcccgtagg acgttcactt ttcttggtgc acagccagcc 180
 cccacagga gcttctctga ggaagttgct gcaacttaacc gctactttgg cggcctgaaa 240
 tctgggtgga atgcttatgt gattggagat ccagcaagac ctggacagaa gtggcacgt 299

<210> 576
 <211> 299
 <212> nucleic acid
 <213> Zea mays

 <400> 576

 ctcatatgtc ctctctgagt caagcttggt tatctatcct ctgaagattg tcatcaagac 60
 ctgtggcaact accaagctcc tgctcaccat tccaagaatc cttgagcttg ctgaagagct 120
 gtctatgcc aattgctgtg tgaagtactc ccgtgggacg ttcactcttc ctggtgcaca 180
 gccagcccc cacaggagct tctctgagga agttgctgca ctttaaccgt actttggcgg 240
 cctgaaatct ggtggtaatg cttatgtgat tggagatcca gcaagacctg gacagaagt 299

<210> 577
 <211> 297
 <212> nucleic acid
 <213> Zea mays
 <400> 577
 gngctcctcc tcctatgaag aaggactaca agctggctaa tttctctgc tgggaggagg 60
 aagcggatgc catggaggag aaggcgggag tgcttgatga gtaagacggg cttctgggggt 120
 cgatttgctt ctgagttgtt tattttatat cgtcgcaatt tcgtggttgt cgtttgggta 180
 ttctgtgaag cagccaagcc aggctattgt tatgaaaatt tgctgtctgt aagcatgtga 240
 acttccgatg ttgccacatg ctggatcagt ctgaataagt aagtatgcag ctctagg 297

<210> 578
 <211> 303
 <212> nucleic acid
 <213> Zea mays
 <400> 578
 aanttctnct acgncaactga gtaccacagag caaccaatgg ttaaccttga gatgtgcatg 60
 actggtctgg acaagaagaa agcttgtgtc tttttcaaga ctaatgctga tgggaacaca 120
 acatgtgcca aggaaatgac aaagctctct ggcattctctg aaatcatccc cgagatggag 180
 atctgogatt ttgacttcga accctgtggc tactccatga atgcgatcca tggctctgca 240
 ttctccacaa tccatgtgac gcccgaggac ggtttcagct acgccagtta cgaggttatg 300
 ggc 303

<210> 579
 <211> 304
 <212> nucleic acid
 <213> Zea mays
 <400> 579
 ccgaggacgg tttcagctat gccagttatg aggttatggg cttggatgcc actgctctgt 60
 cttatggtga ctttgtcaag agggtccttc ggtgctttgg cccctcgag ttttccgttg 120
 ccgtgacat cttcggcggg cgtggccatg ccgggacatg gggaaaggca cttggtgcag 180
 aggtctatga ctgcaacaac atggtggagc aggagctgcc tggaggcggg ctctctgtgt 240

accagagctt ctgtgctgct gaagacgctg ttgctacctc gcccaaactt gttttccact 300
gctt 304

<210> 580
<211> 340
<212> nucleic acid
<213> Zea mays

<400> 580

ttttccactg ctttgacggc gngaacgtgg agagtgtcc tcctcctatg aagaaggact 60
acaaactggc taatcttcta tgctgggagg aggaagcgga tgccatggag gagaaggcgg 120
gagtgttga tgagtaagac gggcttctgg ggtcgatttg cttctgagtt gtttatttat 180
atcgtcgcaa tttcgtgggt gtcgtttggg tatctgtgaa gcagccaagc caagctattg 240
ttatgaaaat tgtcgtctgt aagcatgtga acttccgacg ttgccacatg ctggatcacc 300
tgaataagta agtatgcagc tctaggtgtc agctgcgtct 340

<210> 581
<211> 344
<212> nucleic acid
<213> Zea mays

<400> 581

tggaacaca gcatgtgcca aggaaatgac aaagctctct ggcatctctg aaatcatccc 60
cgagatggag atctgcgatt tgacttgaac cctgtggcta ctccatgaat gcgatccatg 120
gctctgcaat ctccacaatc catgtgacgc ccgaggacgg gttcagccaa gccagnaatg 180
aggctaaggg cctngaagcc actgctctgt cttatggtga ccttgtcaag agggtccttc 240
ggtgctttgg cccctcggag ttttccgttg ccgtgaccat cttcggcggg cgtggccatg 300
ccgggacatg gggaaangca ttggtgcaga ggtctatgac tgca 344

<210> 582
<211> 300
<212> nucleic acid
<213> Zea mays

<400> 582

atctctctgg cctgtgtgct cctgataatc gaccagttcc ccagtcttgc tccttgcaact 60

cctccctcct ccattctccag cattgtgttc tgattcacct gctccaatgg ctgttctttc 120
 tgctgctgat gcttccccgg tctcagctat cgggtttgag ggctatgaga agcgccttga 180
 gatcacattc tctgaggcac ctgtctttgt ggaccctcat gggcgtgggtt tgcgtgccct 240
 ctccagggcc cagattgact ctgttctgga tcttgcaagg tgcacaattg tgtccgagct 300

<210> 583
 <211> 298
 <212> nucleic acid
 <213> Zea mays
 <400> 583

aaggatttcg actcatatgt cctctctgag tcaagcttgt ttatctatcc tctgaagatt 60
 gtcattcaaga cctgtggcac taccaagctc ctgctcacca ttccaagaat ccttgagctt 120
 gctgaagagc tgtctatgcc acttgctgct gtgaagtact cccgtgggac gttcatcttt 180
 cctgggtgcac agccagcccc ccacaggagc ttctctgagg aagttgctgc acttaaccgc 240
 tactttggcg gcctgaaatc tgggtgtaat gcttatgtga ttggagatcc agcaagac 298

<210> 584
 <211> 351
 <212> nucleic acid
 <213> Zea mays
 <400> 584

gcttgotgaa gagctgtcta tgccacttgc tgctgtgaag tactcccgtg ggacgttcat 60
 ctttctctggc gcacagccag cccccccaca ggagcttctc tgaggaagtt gctgcactta 120
 accgetactt tggcggcctg aaatctggtg gtaatgctta tgtgattgga gatccagcaa 180
 gatctggaca gaagtggcac gtcttctacg ccaactgagta ccagagcaa ccaatggtta 240
 accttgagat gtgcatgact ggtctggaca agaagaaagc ttgtgtcttt tcaagactaa 300
 tgetgatngg gaacacaaca tgtgccaag gnaatgacna agctctctng c 351

<210> 585
 <211> 291
 <212> nucleic acid
 <213> Zea mays
 <400> 585

gtgctttggc ccctcagagt tttccgttgc cgtgaccatc ttccggcgggc gtggccatgc 60
 cgggacatgg ggaaaggcac ttggtgcaga ggtctatgac tgcaacaaca tgggtggagca 120
 ggagctgcct ggaggcgggc tcctcgtgta ccagagcttc tgtgctgctg aagacgctgt 180
 tgctacctcg cccaaatctg tttccactg ctttgacggc gagaacgtgg agagtgtctc 240
 tcctcctatg aagaaggact acaagctggc taatcttctc tgctgggagg a 291

<210> 586
 <211> 514
 <212> nucleic acid
 <213> Zea mays

<400> 586

agnnnanatn tgnttttttn gngtgnnnng ananagtaca gcnttaccgg tccggaattc 60
 ccgggtcgac ccacgcgtcc gggagctgcc tggaggcggg ctctcgtgt accagagctt 120
 ctgtgctgct gaagacgctg ntgctacctc gcccaaactc gttttccact gctttgacgg 180
 cgagaacgtg gagagtgtc ctctcctat gaagaaggac tacaagctgg ctaatcttct 240
 ctgctgggag gaggaagcgg atgcatgga ggagaaggcg ggagtgttg atgagtaaga 300
 cgggcttctg gggtcgattt gcttctgagt tgtttatttt atatcgtngc aatttcttgg 360
 ttgtcgttng gttattntgn ggaagnagnn aanccagct atngtaatg aaaanntnnt 420
 tntntaata ntattaactt tnnatnttn tcaanattct tgatnannnt tantaataan 480
 ttatntaatn tntaaggtn nanntttnn ttnt 514

<210> 587
 <211> 299
 <212> nucleic acid
 <213> Zea mays

<400> 587

ggggggcctg aaatctggtg gtaatgctta tgtgattgga gatccagcaa gacctggaca 60
 gaagtggcac gtcttctacg cactgagta ccagagcaa ccaatggtta accttgagat 120
 gtgcatgact ggtctggaca agaagaaagc ttgtgtcttt ttcaagacta atgctgatgg 180
 gaacacaaca tgtgccaagg aaatgacaaa gctctctggc atctctgaaa tcatccccga 240
 gatggagatc tgogattttg acttcgaacc ctgtggctat ccatgaatgc gatccatgg 299

<210> 588
 <211> 288
 <212> nucleic acid
 <213> Zea mays

 <400> 588

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 tacctcgccc aaatctgttt tccactgctt tgacggcgag aacgtggaga gtgctcctcc 120
 toctatgaag aaggactaca agctggctaa tcttctctgc tgggaggagg aagcggatgc 180
 catggaggag aaggcgggag tgcttgatga gtaagacggg cttctggggc cgatttgctt 240
 ctgagttggt tattttatat cgtcgcaatt tcgtggttgc cgtttggt 288

<210> 589
 <211> 305
 <212> nucleic acid
 <213> Zea mays

 <400> 589

 gcttggttat ctatcctctg aagattgtca tcaagacctg tggcactacc aagctcctgc 60
 tcaccattcc aagaatcctt gagcttgctg aagagctgtc tatgccactt gctgctgtga 120
 agtactcccg tgggacgttc atctttcctg gcgcacagcc agccccccac aggagcttct 180
 ctgaggaagt tgctgcactt aaccgctact ttggcggcct gaaatctggt ggtaatgctt 240
 atgtgattgg agatccagca agacctggac agaagtggca gtcttctacg ccaactggagt 300
 accca 305

<210> 590
 <211> 295
 <212> nucleic acid
 <213> Zea mays

 <400> 590

 gggtccttcg gtgctttggc cctcggagt tttccgttgc cgtgaccatc ttcggcgggc 60
 gtggccatgc cgggacatgg ggaaaggcac ttggtgcaga ggtctatgac tgcaacaaca 120
 ttggtggagca ggagctgcct ggaggcgggc tctcgtgta ccagagcttc tgtgctgctg 180
 aagacgctgt tgctacctcg cccaaatctg ttctccactg ctttgacggc gagaacgtgg 240

agagtgtcc tctcctatg aagaaggact acaaactggc taatcttcta tgctg 295

<210> 591
 <211> 298
 <212> nucleic acid
 <213> Zea mays
 <400> 591

gcgggagtgc ttgatgagta agacgggctt ctgggggtoga tttgcttctg agttgtttat 60
 tttatatcgt cgcaatttcg tggttgtcgt ttggttatto tgtgaagcag ccaagccagg 120
 ctattgttat gaaaatttgt cgtctgtaag catgtgaact tccgatgttg ccacatgctg 180
 gatcagtctg aataagtaag tatgcagctc taggtggtea gctgcgtcta ccacaatgag 240
 catgaacgta tggagaaata tctgtgaacc catttgttta tgaataagat tgtttttt 298

<210> 592
 <211> 320
 <212> nucleic acid
 <213> Zea mays
 <400> 592

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 cccccaccgg agcttctccg aggaagttgc tgtacttaac cgatactttg ggggcctgaa 120
 gtctggtggc aatgcttatg tgattggaga tgcagcaaga ccaggacaga agtggcacat 180
 ctactacgcc actgagtacc cagagcaacc aatgggtcaac cttgagatgt gcatgactgg 240
 tctggacacg aagaaagctt cagtcttctt caagactaat gctgatggca acacaacatg 300
 tgccaaggaa atgacgaagt 320

<210> 593
 <211> 284
 <212> nucleic acid
 <213> Zea mays
 <400> 593

tggctactcc atgaatgcga tccatggctc tgcattctcc acaatccatg tgacgcccga 60
 ggacggtttc agctacgcca gttacgaggt tatgggcttg gatgccactg ctctgtctta 120
 tgggtgacctt gtcaagaggg tctccgggtg ctttggcccc tcagagtttt ccgttgccgt 180

gaccatcttc ggcgggcgtg gccatgccg gacatggga aaggcacttg gtgcagaggt 240
ctatgactgc aacaacatgg tggagcagga gctgcctgga ggcg 284

<210> 594
<211> 285
<212> nucleic acid
<213> Zea mays

<400> 594

gaagtccag tctgctgctt actccaactg cgcaagaag ccatcctgat atcccttttg 60
gcttcctcat tctagtagtt taggatttct tttctgacac tttgattctg accaatctct 120
ctggcctgct gcttcctgat aatcgaccag ttccccagtc ttgctccttg cactcctccc 180
tccatctcca gcattgtggt ctgattcacc tgcaccaatg gctgtttctt ctgctgctga 240
tgcttccccg gtctcagcta tcgggtttga ggcctatgag aagcg 285

<210> 595
<211> 286
<212> nucleic acid
<213> Zea mays

<400> 595

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tctgaccaat ctctctggcc tgctgcttcc tgataatcga ccagttcccc agtcttgctc 240
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<213> Zea mays

<400> 596

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agtttaggat ttcttttctg acactttgat tctgaccaat ctctctggcc tgctgcttcc 180

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 <213> Zea mays
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<210> 602
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 <212> nucleic acid
 <213> Zea mays
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 <212> nucleic acid
 <213> Zea mays
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<400> 606

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<400> 607

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<400> 608

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<400> 611

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<400> 612

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 <212> nucleic acid
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<400> 613

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<400> 614

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<400> 615

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<400> 616

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<400> 617

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 <212> nucleic acid
 <213> Zea mays

<400> 618

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 <212> nucleic acid
 <213> Zea mays

<400> 619

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 aagctcctgc tcaccattcc aagaatcctt gagcttgctg aagagctgtc tatgccactt 180

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<210> 620
<211> 275
<212> nucleic acid
<213> Zea mays

<400> 620

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<210> 621
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<212> nucleic acid
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<400> 621

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<210> 622
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<212> nucleic acid
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<400> 622

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<210> 623
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<212> nucleic acid
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<400> 623

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<210> 624
<211> 402
<212> nucleic acid
<213> Zea mays

<400> 624

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<213> Zea mays

<400> 625

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 <213> Zea mays
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<213> Zea mays
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<211> 263
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 <400> 633

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 gtttgcgtnc cctctccagg gccagattg nctntgttct ggatcttgca cgggtgcacaa 180
 tgtgtctgag ctctccaaca aggatttcga ctcatatgtc ctttctgagt caagcttggt 240
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<210> 634

<211> 292
 <212> nucleic acid
 <213> Zea mays

 <400> 634

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 atctttcctg gcgcacagcc agccccccac aggagcttct ctgaggaagt tgctgcactt 180
 aaccgctact ttggcggcct gaaatctggt ggtantgctt atgtgattgg agatccagca 240
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<210> 635
 <211> 270
 <212> nucleic acid
 <213> Zea mays

 <400> 635

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 ctgctgctta ctccaactgc gcgaagaagc catcctgata tcccttttgg ctctctcatt 180
 ctagtagttt aggatttctt ttctgacact ttgattctga ccaatctctc tggcctgctg 240
 ctctctgata atcgaccagt tccccagtct 270

<210> 636
 <211> 270
 <212> nucleic acid
 <213> Zea mays

 <400> 636

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 aagttccagt ctgctgctta ctccaactgc gcgaagaagc catcctgata tcccttttgg 120
 ctctctcatt ctagtagttt aggatttctt ttctgacact ttgattctga ccaatctctc 180
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 ccatctccag cattgtgttc tgattcacct 270

<210> 637

<211> 267
 <212> nucleic acid
 <213> Zea mays

 <400> 637

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 gctatgagaa ggccttgag atcacattct ctgaggcacc tgtctttgtg gaccctcatg 180
 ggcgtgggtt gcgtagcctc tccagggccc agattgactc tgttctggat cttgcacggt 240
 gcacaattgt gtccgagctc tccaaca 267

<210> 638
 <211> 281
 <212> nucleic acid
 <213> Zea mays

 <400> 638

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 aggactacaa gctggctaata cttctctgct gggaggagga agcggatgcc atggaggaga 180
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 attttatata gtgcgaattc gtggntgtcg ttggttattc t 281

<210> 639
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 639

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 gtctttttca agactaatgc tgatgggaac acaacatgtg ccaaggaaat gacaaagctc 180
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 ggctactcca tgaatgcga 259

<210> 640

<211> 298
 <212> nucleic acid
 <213> Zea mays

 <400> 640

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 gagctctcca acaaggattt tgacncctat gttctctccg agtcaagctt gttcatctat 180
 cctctgaaga ttgtcatcaa gacctgtggc actaccaagc tcctgctcac aattccaagg 240
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<210> 641
 <211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 641

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 acottgagat gtgcatgact ggtctggaca agaagaaagc ttctgtcttt tnccaagact 180
 aatgctgatg ggaacacagc atgtgccaaag gaaatgacaa agctctctgg catctctgaa 240
 atcatccccg agatggagat ctgcgatttt g 271

<210> 642
 <211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 642

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 acagagctgt ctatgccact tgctgctgtg aagtactccc gtgggacgtt catctttcct 120
 ggtgcacagc cagcccccca caggagcttc tctgaggaag ttgctgcact taaccgctac 180
 tttggcggcc tgaaatctgg tggtaatgct tatgtgattg gagatccagc aagacctgga 240
 cagaagtggc acgtcttcta cgccactgag t 271

<210> 643

<211> 273
 <212> nucleic acid
 <213> Zea mays

 <400> 643

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 atgggggaaag gcacttggtg cagaggtcta tgactgcaac aacatgggtg agcaggagct 120
 gcctggaggc gggctontcg tgtaccagag cttctgtgct gctgaagacg ctgttgctac 180
 ctcgccccaa tctgttttcc actgctttga cggcgagaac gtggagagtg ctcctcctcc 240
 tatgaagaag gactacaagc tggctaattc tct 273

<210> 644
 <211> 257
 <212> nucleic acid
 <213> Zea mays

 <400> 644

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 gggacatggg gaaaggcact tgggtgcagag gtctatgact gcaacaacat ggtggagcag 180
 gagctgctg gaggggggct cctcgtgtac cagagcttct gtgctgctga agacgtgtt 240
 gctacctcgc ccaaatc 257

<210> 645
 <211> 310
 <212> nucleic acid
 <213> Zea mays

 <400> 645

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 tctgagtcaa gcttgtttat ctatcctctg aagattgtca tcaagacctg tggcactacc 120
 aagctcctgc tcaccattcc aagaatcctt gagcttgctg aagagctgtc tatgccactt 180
 gctgctgtga agtactcccg tgggacgttc atctttcctg gtgcacagcc agccccccac 240
 aggagcttct ctgaggaagt tgctgcactt aaccgctatt tggggcctga aatctgggtg 300
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<210> 646
 <211> 278
 <212> nucleic acid
 <213> Zea mays

 <400> 646

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 ccaggctatt gttatgaaaa tttgtcgtct gtaagcatgt gaacttccga tgttgccaca 180
 tgctggatca gtctgaataa gtaantatgc agctctaggt ggtcagctgc gtctaccaca 240
 atgagcannng acntnntgga gaaatatctg tgaacccc 278

<210> 647
 <211> 257
 <212> nucleic acid
 <213> Zea mays

 <400> 647

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 cctcctatga agaaggacta caagctgggt aatcttctct gctgggagga ggaagcggat 180
 gccatggagg agaaggcggg agtgcttgat gagtaagacg ggcttctggg gtcgatttgc 240
 ttctgagttg ttaattt 257

<210> 648
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 648

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 actacaaact ggctaattct ctatgctggg aggaggaagc ggatgccatg gaggagaagg 180
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<210> 649
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 649

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 tgggtggagca ggagctgcct ggaggcgggc tctcgtgta ccagagcttc tgtgctgctg 180
 aagacgctgt tgctacctcg cccaaatctg ttttccactg ctttgacggc gagaacgtgg 240
 agagtgtctc tctctctat 259

<210> 650
 <211> 256
 <212> nucleic acid
 <213> Zea mays

 <400> 650

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 gggacatggg gaaaggcact tgggtgcagag gtctatgact gcaacaacat ggtggagcag 180
 gagctgcctg gaggcgggct cctcgtgtac cagagcttct gtgctgctga agacgctggt 240
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<210> 651
 <211> 267
 <212> nucleic acid
 <213> Zea mays

 <400> 651

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 atccatggct ctgcattctc cacaatccat gtgacgccg aggacggttt cagctatgcc 180
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 gtccttcggt gctttggccc ctcgag 267

<210> 652
 <211> 531
 <212> nucleic acid
 <213> Zea mays
 <400> 652
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 agaggtgccg ggaccgagga tttggccgag ttctaggtag agaatcgggtt ttacttgtct 180
 cagccccggg tctgctgcgt ctggtggtgg tgaaggggag aaattcgtga gatctgttcc 240
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 tgatgtacta atggagtcta aggggtggcaa aaagtctagc agtagtcgtt ctatgatgta 360
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 tccagtctgc tgcttactcc aactgcgcga agaagccatc ctgatatccc ttttggttcc 480
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<210> 653
 <211> 279
 <212> nucleic acid
 <213> Zea mays
 <400> 653
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 gcattgtggt ctgattcacc tgctccaatg gctgttctt 279

<210> 654
 <211> 488
 <212> nucleic acid
 <213> Zea mays
 <400> 654
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cctnttagcg tggccatgcc gggacatggg gaaaggcact tgggtgcagag gtctatgatg 120
 caacaacatg gtggagcang agctgcctgg aggcgggctc ctcntgtacc aaagctttgt 180
 nctgntgaag acgcngttgc tanctcgccc aaatccgttt tccantgntt tgacnggaaa 240
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 ctggggncna ttttcgtctn agttngtann ttttaagcnt cccaanattc cccgggtgtc 420
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 nccgnnng 488

<210> 655
 <211> 252
 <212> nucleic acid
 <213> Zea mays

<400> 655
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 cttgtttatc tatcctctga agattgtcat caagacctgt ggccactacca agctcctgct 180
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 gtactcccgt gg 252

<210> 656
 <211> 282
 <212> nucleic acid
 <213> Zea mays

<400> 656
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 gagatgcagc aagaccagga cagaagtggc acatctacta cgccactgag taccagagc 180
 aaccaatggt caaccttgag atgtgcatga ctggctctga cacgaagaaa gcttcagtct 240
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<210> 657

<211> 250
 <212> nucleic acid
 <213> Zea mays

 <400> 657

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 ggcgcacagc cagcccccca caggagcttc tctgaggaag ttgctgcact taaccgctac 180
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<210> 658
 <211> 261
 <212> nucleic acid
 <213> Zea mays

 <400> 658

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 atgtcctttc tgagtcaagc ttgtttatct atcctctgaa gattgtcatc aagacctgtg 180
 gcactaccaa gctcctgctc accattccaa gaatccttga gcttgctgaa gagctgtcta 240
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<210> 659
 <211> 261
 <212> nucleic acid
 <213> Zea mays

 <400> 659

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 actaatgctg atgggaacac aacatgtgcc aaggaaatga caaagctctc tggcatctct 180
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<210> 660

<211> 454
 <212> nucleic acid
 <213> Zea mays

<400> 660

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gcgtctggtg gtggtgaagg ggagaaattc gtgagatctg ttccggatca ggcgtgctgag   180
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tctaaggggtg gcaaaaagtc tagcagtagt cgttccatga tgtatgaagc tccccttggc   300
tacagcattg aggacgttcg acctgccgga ggcgtgaaga agttccagtc tgctgcttac   360
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<210> 661
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 <212> nucleic acid
 <213> Zea mays

<400> 661

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cacgtcttct acgccactga gtaccagag caaccaatgg ttaaccttga gatgtgcatg   180
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<210> 662
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 <212> nucleic acid
 <213> Zea mays

<400> 662

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atgaggttat gggcttggat gccactgccc tgtcttacgg tgaccttgtc aagagggtcc   180

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tctgtgctgc tga 253

<210> 666
<211> 248
<212> nucleic acid
<213> Zea mays

<400> 666

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gtaatgctta tgtgattgga gatccagcaa gacctggaca gaagtggcac gtcttctacg 180
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agaagaaa 248

<210> 667
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 667

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<210> 668
<211> 247
<212> nucleic acid
<213> Zea mays

<400> 668

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gaaagcttgt gtctttttca agactaatgc tgatgggaac acaacatgtg ccaaggaaat 180

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<210> 669
<211> 445
<212> nucleic acid
<213> Zea mays

<400> 669

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ctgacacttt gattctgacc aatct 445

<210> 670
<211> 276
<212> nucleic acid
<213> Zea mays

<400> 670

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<210> 671
<211> 283
<212> nucleic acid
<213> Zea mays

<400> 671

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 ccaactgogc aaagaagcca tcttgatata ccttttggnt tcctcgttct agtagtttag 180
 gatttttttt ctgacacttc gnttcttacc aatccccctg gcctgctgct ttcttgacaa 240
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<210> 672
 <211> 284
 <212> nucleic acid
 <213> Zea mays

<400> 672

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 gctccaatgg ctgttctttc tgctgctgat gttccccggt ctgagctatc gggtttgagg 180
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<210> 673
 <211> 443
 <212> nucleic acid
 <213> Zea mays

<400> 673

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<210> 674

<211> 254
 <212> nucleic acid
 <213> Zea mays

<400> 674

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 tcgtgtacca gagcttctgt gctgctgaag acgctgttgc tacctcgccc aaatctgttt 180
 tccactgctt tgacggcgag aacgtggaga gtgctcctcc tcctatgaag aaggatacaa 240
 gctggctaatt cttc 254

<210> 675
 <211> 272
 <212> nucleic acid
 <213> Zea mays

<400> 675

ctgaagtctg gtggcaatgc ttatgtgatt ggagatgcag caagaccagg acagaagtgg 60
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 actggtctgg acacgaagaa agcttcagtc ttcttcaaga ctaatgctga tggcaacaca 180
 acatgtgcc aaggaaatgac gaagctctct ggtatctctg aaattatccc tgagatggag 240
 atctgtgatt ttgacttcga accctgcggc ta 272

<210> 676
 <211> 240
 <212> nucleic acid
 <213> Zea mays

<400> 676

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 tcaagactaa tgctgatggg aacacaacat gtgccaaagga aatgacaaag ctctctggca 180
 tctctgaaat catccccgag atggagatct gcgattttga cttcgaacct tgtggctact 240

<210> 677
 <211> 243
 <212> nucleic acid

<213> Zea mays

<400> 677

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ggtaatgctt atgtgattgg agatccagca agacctggac agaagtggca cgtcttctac 180
gccactgagt acccagagca accaatgggt aaccttgaga tgtgcatgac tgggtctggac 240
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<210> 678

<211> 243

<212> nucleic acid

<213> Zea mays

<400> 678

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tatgactgca acaacatggg ggagcaggag ctgcctggag gcgggctcct cgtgtaccag 180
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gac 243

<210> 679

<211> 251

<212> nucleic acid

<213> Zea mays

<400> 679

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ggacgttcga cctgccggag ggtgaagaag ttccagtctg ctgcttactc caactgcgcg 180
aagaagccat cctgatatcc cttttggcct cctcattcta gtagtttagg atttcttttc 240
tgacactttg a 251

<210> 680

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 680

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cgaggaagtt gctgtactta accgatactt tggggggcctg aagtctgggtg gcaatgctta 180
tgtgattgga gatgcagcaa gaccaggaca gaagtggcac atctactacg ccactgagta 240
cccagagcaa ccaatggtca accttgag 268

<210> 681

<211> 249

<212> nucleic acid

<213> Zea mays

<400> 681

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atccatggct ctgcattctc cacaatccat gtgacgccg aggacggttt cagcnatgcc 180
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gtccttcgg 249

<210> 682

<211> 241

<212> nucleic acid

<213> Zea mays

<400> 682

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atctgggtgg aatgcttatg tgattggaga tccagcaaga cctggacaga agtggcacgt 180
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<210> 683

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 683

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tggctactcc atgaatgcga tccatggctc tgcattctcc acaatccatg tgacgcccga 180
ggacgggtttc agctacgccg gttacgaggt tatgggctnt anttgccatg ctctgtctta 240
tggtgacctt gtcaagaggt cctccggt 268

<210> 684

<211> 264

<212> nucleic acid

<213> Zea mays

<400> 684

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ctacaaaacta gctaattctt tctgtctgnnn ggaggaagcg gatgccatgg aggagaaggc 120
gggagtgttt gatgagtaag acgggcttct ggtgtcgatt tgcttctgag ttgtttatTT 180
tatatcgtcg cnatttcgtg gttgtcgttt ggttatctgt gaagcagcca agccaggcta 240
ttgttatgaa aattgtcgtc tgta 264

<210> 685

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 685

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aagactaatg ctgatgggaa cacaacatgt gccaaaggant ganaancctc tctggcatct 180
ctgaaatcat ccccgagatg gagatctgcg attttgactt cgaaccctgt ggctactcca 240
tgaatgcgat ccatggctct gcattctc 268

<210> 686

<211> 332

<212> nucleic acid

<213> Zea mays
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 gggagtgctt gatgagtaan acgggcttct ggggtcgatt tgcttctgag ttgtttatct 180
 tatatcgctg caatttcgtg gttgtcgctt ngttattctg tgaagcagcc aagccaagct 240
 attgttatga nntttgtcgt ctgtaagcat gtgacttccg ccgttgccac atgctggatc 300
 agtctgaata gtaagtngca gctctagtgg nc 332

<210> 687
 <211> 237
 <212> nucleic acid
 <213> Zea mays

<400> 687
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 tggatttgc tctgagttgt ntattttata tgcgcgcaat ttcgtgggtg tggtttggtt 120
 attctgtgaa gcagccaagc caggctattg ttatgaaaat ttgtcgtctg taagcatgtg 180
 aaattccgat gttgccacat gctggatcag tctgaataag taagtatgca gctctag 237

<210> 688
 <211> 286
 <212> nucleic acid
 <213> Zea mays

<400> 688
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 ccgggtctca gctatcgggt ttgagggtta tgagaagcgc cttgagatca cattctctga 120
 ggcacctgtc ttgttggaac ctcatggcgt gggttgctg cctctccag ggcccagatt 180
 gactctgttc tggatcttgc acggtgcaca atgtgtccga gctctccaac aaggatttcg 240
 actcatagtc ctctctgagt caagcttggt tatctatctc tgaaga 286

<210> 689
 <211> 300
 <212> nucleic acid

<213> Zea mays

<400> 689

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actaatggag tctaagggtg gcaaaaagtc tagcagtagt cgttctatga tgtatgangc 120
tccccttggc tacagcnttg aggacgttcg acctgccga cggatgaagaa gttccagtct 180
gctgcttact ccaagtgcgc gangaagcca tcttgatc cttttggctt cctcattct 240
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<210> 690

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 690

gtgattggag gccgcaaag tagcggntaa gtgcagcaac ttcctnagag aagctcctgt 60
ggggggctgg ctgtgcacca ggaaagatga acgtcccang ggagtacttc acagcagcna 120
gtggcataga cagctcttca gcaagctcaa ggattcttgg aatggtgagc aggacttggt 180
agtgccacag gtcttgatga caatcttcag aggatagatn aacaagcttg actcatagag 240
gacatatgat ccgaaatcct tgtggagagc tcgacaca 278

<210> 691

<211> 525

<212> nucleic acid

<213> Zea mays

<400> 691

gnnaantgtt ttgtaanggg ggggnnaang nnaagctttt accggtccgg aattcccggg 60
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ctgagagggtg ccgggaccga ggaactggcc gactactagg tagagaatcg gtttttcttg 180
tctcagcccg ggggtctgtg cgtctgggtg tggatgaagg gagaaattcg tgagatctgt 240
tccggatcag ggtgagagc tcgggaatca ggggtttcac acatagcttc gtcgatttga 300
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gtatgaagct ccccttggct acagcattga ggacgttcga cctgccggag gctgaagaa 420

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 <212> nucleic acid
 <213> Zea mays
 <400> 692

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 tggagatctg cgatttttgac ttcgaaccct gtggctactc catgaatgcg atccatggct 180
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 ta 242

<210> 693
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 gtggtagacg cagctgacca cctagagctg catacttact tattcagact gatccagcat 120
 gtggcaacgt cggaagttca catgcttaca gacgacaaat tttcataaca atagcttggc 180
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<210> 694
 <211> 263
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 <400> 694

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 cttganatgn gcatgactgg tctggacnag nagaangctt gtgtcttttt caagantaat 180
 gctgatgggn acacnacatg tgccaaggga atgacnaagc tcnctggcat ctngaaagc 240

ntnccccgnga tggannntgc gnt

263

<210> 695
<211> 245
<212> nucleic acid
<213> Zea mays

<400> 695

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aggcacttgg tgcagaggtc tatgactgca acaacatggt ggagcaggag ctgcctggag 180
gogggtect cgtgtaccag agcttctgtg ctgctgaaga cgctgttgc actcgcccaa 240
atctg 245

<210> 696
<211> 229
<212> nucleic acid
<213> Zea mays

<400> 696

gctgctgatg cttccccggt ctcagctatc gggtttgagg gctatgagaa gcgccttgag 60
atcacattct ctgaggcacc tgtctttgtg gacctcatg ggctgggttt gcgtgccctc 120
tccaggcccc agattgactc tgttctggat cttgcacggt gcacaattgt gtctgagctc 180
tccaacaagg atttcgactc atatgtcctt tctgagtcaa gcttgttta 229

<210> 697
<211> 228
<212> nucleic acid
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<400> 697

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actgcaaaa catggtggag caggagctgc ctggaggcgg gtcctcgtg taccagagct 120
tctgtgctgc tgaagacgt gttgctacct cgcccaaadc tgtttccac tgctttgacg 180
gcgagaacgt ggagagtgc ctcctccta tgaagaagga ctacangc 228

<210> 698

<211> 227
<212> nucleic acid
<213> Zea mays

<400> 698

caacatggtg gagcaggagc tgccctggagg cgggctcctc gtgtaccaga gcttctgtgc 60
tgctgaagac gctgttgcta cctcgcccaa atctgttttc cactgctttg acggcgagaa 120
cgtggagagt gctcctcctc ctatgaagaa ggactacaag ctggctaate ttctctgctg 180
ggaggaggaa gcggatgcca tggaggagaa ggcgggagtg cttgatg 227

<210> 699
<211> 275
<212> nucleic acid
<213> Zea mays

<400> 699

gtgattttga cttcgaaccc tgcggctact ccatgaatgc aatccatggc tctgcgttct 60
ccacgatcca tgtgacgcct gaggatgggt tcagctacgc cagttatgag gttatgggct 120
tggatgccac tgccctgtct tacggtgacc ttgtcaagag ggtccttggg tgcttcggcc 180
cctcggaatt ttctgtcgcc gtgaccatct tcggcggggc gggccaagct gggacatggg 240
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<210> 700
<211> 497
<212> nucleic acid
<213> Zea mays

<400> 700

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ctgagaggtg ccgggaccga ggaactggcc gagttctagg tagagaatcg gttttgcttg 180
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tccggatcag gcgtgcgagc tcgggaatca ggggtttcac acatagcttc gtcgatttga 300
atattgatga ctaatggagt ctaagggttg caaaaagtct agcagtagtc gttccatgat 360
gtatgaagct ccccttggtc acagcattga ggacgttcga cctgccggag gcgtgaagaa 420

gttccagtct gctgcttact ccaactgcgc gaagaagcca tcttgatata ccttttggct 480
tcttcattct agtagtt 497

<210> 701
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<212> nucleic acid
<213> Zea mays

<400> 701

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tacagcattg aggacgttcg acctgccgga ggggaagaa gttccagtct gctgcttact 180
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<400> 702

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tccccttggc tacagcattg aggacgttcg acctgccgga ggggaagaa anttccagtc 180
tgctgcttac tccaactgcg cgaagaagcc atcctgatan ccttttnggc ttctcattn 240
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<210> 703
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<400> 703

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<210> 704
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<400> 704

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 gtttgagggc tatgagaagc gccttgagat cacattctct ganncacctg tctttgtgga 180
 ccccatggc agcgggttgc gtgccctctc caggctccag attgactctg ttctggatct 240
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<400> 705

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 tgcaacttaac cgtactttg gcgggctgaa atctgggtgt aatgcttatg tgattggaga 180
 tccagcaaga acctggacag aagtgnacg tcttctaacg caactgagta ccnagagcaa 240
 ccaatggtta ccttgag 257

<210> 706
 <211> 221
 <212> nucleic acid
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<400> 706

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 tccagtctgc tgcttactcc aactgcgcga agaagccatc ctgatatccc ttttggcttc 120
 ctcattctag tagtttagga tttcttttct gacactttga ttctgaccaa tctctctggc 180
 ctgctgcttc ctgataatcg accagttccc cagtcttgc c 221

<210> 707
 <211> 534
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 <400> 707

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 gtctccctga gaggtgccgg gancgaggaa ctggccgagt tctaggtaga gaatcggttt 180
 tgcttgctc agcacggggt ctgctgcgtc tgggtggtggt gaaggggaga aattcgtgag 240
 atctgttccg gatcaggcgt gcgagctcgg gaatcagggg tttcacacat agcttcgtcg 300
 atttgaattt gatgtactaa tggagtctaa ggggtggcaaa aagtctagca gtagtcgttc 360
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 gaagaagttc cagtcttgtt tgcttactcc aactgcgcga agaagccatc ctgatatccc 480
 ttttggtttc ctaatttnan nanttttagga attntttttt gaaaccttnn annt 534

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 <213> Zea mays

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 attcacctgc tccaatggct gttctttctg ctgctgatgc ttccccggtc tcagctatcg 180
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ccagttcccc agtcttgctc cctgcagtcc tccctcctcc atctccagcg ttgtgttctg 240
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ctttgggggc ctgaagtctg gtggcaatgc ttatgtgatt gnagatgcag caagaccagg 180
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tggatgccac tgccctgtct tacggtgacc ttgtcaagag ggtccttggg tgcttcggcc 180
cctcggaatt ttctgtcgcc gtgaccatct tcggcgggcg gggccaagct gggacatggg 240
gaaaggaatt gg 252

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ttgagggtta tgagaagcgc cttgagatca cattctctga ggacacctgtc tttgtggacc 180
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cacggtgcac gatcgtgtcc gag 263

<210> 713
<211> 266
<212> nucleic acid
<213> Zea mays

<400> 713

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caggctattg ttatgaaaat tgcgtctctgt aagcatgtga acttccgatg ttgccacatg 180
ctggancatc tgaataagta agtangnngc tctaggtagt cagctgcgtc tccccacaat 240
gagcatgaag tatggagaaa tatctg 266

<210> 714
<211> 242
<212> nucleic acid
<213> Zea mays

<400> 714

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cttggatgcc actgctctgt cttatgggtga ccttgtcaag agggtcctcg gtgctttggc 180
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ga 242

<210> 715
<211> 257
<212> nucleic acid
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<400> 715

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gttctggatc ttgcacgggtg cacgatcgtg tccgagctct ccaacaagga ttttgantcc 180
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ggcataccaa gctcctg 257

<210> 716
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<212> nucleic acid
<213> Zea mays
<400> 716

ctcagctatc gggtttgagg gctatgagaa gcgccttgag atcacattct ctgaggnacc 60
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tgttcnggat cttgcanggt gnacaattgn gnccgagctc tccaacaagg atttcgactc 180
natgtcctct ctgagtcaan ttgn ttatct atcctctgaa gattgtcatc aagacctgtg 240
gcatancaag ctctgtcan catcnaagaa tcttgagctg ctgaagagct gtctagccat 300
tgtctnctgtg antatccgtg ggagtc 326

<210> 717
<211> 496
<212> nucleic acid
<213> Zea mays
<400> 717

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gagaggtgcc gggaccgagg atttggccga gttctaggtg gagaatcggg tttacttgtc 180
tcagcccggg gtctgctgcg tctgggtgtg gtgaagggga gaaattcgtg agatctgttc 240
cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt cgatttgaat 300
ttgatgtact aatggagtct aagggtggca aaaagtctag cagtagtcgt tctatgatgt 360
atgaagctcc ccttggttac agcattgagg acgttcgacc tgccggaggc gtgaagaagt 420
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cattctaagt anttta

496

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<400> 718

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tgaataagta agtatgcagc tctaggtggt cagctgcgtc taccacaatg agcatgtacg 180
tatggagaaa tatctgtgaa cccattttgt ttatgaataa gatttgtttt ttogagt 237

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<210> 719
<211> 493
<212> nucleic acid
<213> Zea mays

<400> 719

acctnctccn gaattccggg gtcgacccan gcgtgcgcgg gacccgagga actggccgag 60
ttctaggtag agaatcggtt ttgcttgtct cagcacggng tgtgngngt ctggtggtgg 120
ngaaggggag aaattcgtga gatctgttcc ggatcaggcg tncgagctcg ggaatcaggg 180
gtttcacaca tagcttcgtc gatttgaatt tgatgtacta atggagtcta aggggtggcaa 240
aaagtctagc agtagtcgtt ccatgatgta tgaagctccc cttggctaca gcattgagga 300
cgttcgacct gccggaggcg tgaagaantt ccagtctgct gcttactcca actgcgcgaa 360
aaaaccatcc tgatatccct ttggccttcc ctcaatctaa tantttaagg atttcttttt 420
ttgacacttt taattcttac caaatntttt ttggcctggn tgctttcctg anaaatngac 480
canttcccca nnt 493

<210> 720
<211> 264
<212> nucleic acid
<213> Zea mays

<400> 720

tgctttgacg gcgagaacgt ggagagtgct cctcctccta tgaagaagga ctacaaactg 60

gctaattcttc tctgctggga ggaggaagcg gatgccatgg aggagaaggc gggagtgcctt 120
gatgagtaag acgggcttct ggtgtcgatt tgcttctgag ttgtttatatt tatatcgctcg 180
caatttcgtg gttgtcgttt ggttatccgt gaagcaccca gccagcnaat gttagnaaat 240
ttgccgccgt aaccagggaa cttc 264

<210> 721
<211> 272
<212> nucleic acid
<213> Zea mays
<400> 721

caggcgtgcg agctcgggaa tcaggggttt cacacatagc ttcgctcgatt tgaatttgat 60
gtactaatgg agtctaaggg tggcaaaaag tctagcagta gtognntcca tgatgtatga 120
agctccccctt ggctacagca ttgaggacgt tcgacctgcc ggagggtgaa agangttcca 180
gtengengct tactccaact gcnnaagaa gccattcct gatatccctt tnggcttcen 240
cattctagta ggtaggatt ncttttctna ca 272

<210> 722
<211> 487
<212> nucleic acid
<213> Zea mays
<400> 722

ttatntncca ntgnccgnct caaatacana gncncgtntt tcattacaan ncnnnnnnnn 60
nnnnnnnnnn nnncaaagaa ttccctcccc tagccgccgc cgccgctcgt ctccctgaga 120
ggtgccggga ccgaggaact ggccgagttc taggtagaga atcggttttg cttgtctcag 180
cacgggggtct gctgcgtctg gtggtggtga aggggagaaa ttcgtgagat ctggtccgga 240
tcaggcgtgc gagctcggga atcaggggtt tcacacatag cttcgtcgat ttgaatttga 300
tgtactaatg gagtctaagg gtggcaaaaa gtctagcagt agtcgttcca tgatgtatga 360
agctccccctt ggctacagca ttgaggacgt tcgacctgcc gggaggcgtg aagaagttcc 420
agtctgctgc ttactccaac tgcgcgaaag aagccatcct gatatccctt ttggcttcct 480
caatcta 487

<210> 723

<211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 723

 tttgttgatt tgactatgga acataaggng gagatgaagg ttgtggtcag gacagttatt 60
 tcactcacca aggaggaagc g gatgccatg gaggagaagg cgggagtgtc tgatgagtaa 120
 gacgggcttc tgggtgtcgat ttgcttctga gttgtttatt ttatatcgtc gcaatttcgt 180
 gggttgcgtt tgggttattct gtgaagcagc caagccaggc tattgttatg aaaatttcgtc 240
 gtctgtaagc atgtgaactt ccgatgttgc c 271

<210> 724
 <211> 214
 <212> nucleic acid
 <213> Zea mays

 <400> 724

 ctctctggca tctctgaaat catccccgag atggagatct gcgattttga ctttgaaccc 60
 tgtggctact ccataaatgc gatccatggc tctgcattct ccacaatcca tgtgacgccc 120
 gaggaacggt tcagctatgc cagttatgag gttatgggct tggatgccac tgctctgtct 180
 tatggtgacc ttgtcaagag ggtcttcggt gttg 214

<210> 725
 <211> 260
 <212> nucleic acid
 <213> Zea mays

 <400> 725

 caatcgacca gttccccagt cttgctccct gcagtcctcc ctctccatc tccagcgttg 60
 tgttctgact cacctgcacc aatggctgtt ctttctgctg ctgggtgctc cccggcctca 120
 gctatcgggt ttgagggcta tgagaagcgc cttgagatca cattctctga ggcacctgtc 180
 tttgtggacc cccatggcag cggtttgcgt gccctctcca ggtcccagat tgactctgtt 240
 ctggatcttg cacggtgcac 260

<210> 726
 <211> 536
 <212> nucleic acid

<213> Zea mays

<400> 726

gagaaantgg cttccttnta gnagnnnnncnn angganagnn ttctatcctn tcnantttcc 60
 cgnatgngac cgacgcgtcc gcaangaatt ccctccccta gccgcgcgt cgcgtcgctcg 120
 tctccctgag aggtgccggg ncnggggaac tggccgagta ctaggtagag aatcggtttt 180
 tcttgtctca gcccggggtc tgctgcgtct ggtggtggtg aaggggagaa attcgtgaga 240
 tctgttccgg atcaggcgtg cgagctcggg aatcangggg ttcacacata gcttcgtcga 300
 tttgaatttg atgtactaat ggagtctaag ggtggcaaaa agtctagcag tagtcgttct 360
 atgatgtatg aagctnccct tggctacagc attgaggacc gttcgacctg ccggangcgt 420
 gaagaagttc cagtctnctg cttacttcaa ctgccnaag aagccatcct natannccct 480
 ttggctttct tattctaata gntttangat ttctttttct ganaccnnt attttt 536

<210> 727

<211> 456

<212> nucleic acid

<213> Zea mays

<400> 727

caaagnatlc nctcccctag ccgcgcgcgc cgctcgnctc cctgagaggt gccgggaccg 60
 aggatltggc cgagttctag gtagagaatc ggttttactt gtctcagccc ggggtctgct 120
 gcgtctggtg gtggtgaagg ggagaaattc gtgagatctg ttccggatca ggcgtagcag 180
 ctcggaatc aggggtttca cacatagctt cgtcgatttg aatttgatgt actaatggag 240
 tctaagggcg gcaaaaagtc tagcagtagt cgttctatga tgtatgaagc tccccttggc 300
 tacagcattg angacgttcg acctgccnga ngcgtgaaga aattncaagt ctgctgctta 360
 ctccaactgc gccaaaaaac catnctgata ttcttttggg ctttcctaata ctagtaagtt 420
 aaggatttct tttctgacac ttttgaatct gancca 456

<210> 728

<211> 196

<212> nucleic acid

<213> Zea mays

<400> 728

ggtgtcgatt tgcttctgag ttgtttatTT tatatcgTcg caatttTcgTg gttgtcgTTt 60
 ggTtattctg tgaagcagcc aagccaggct attgttatga aaatttTgTcg tctgtaagca 120
 tgtgaacttc cgatgttgcc acatgctgga tcagtctgaa taagtaagta tgcagctcta 180
 ggtggTcagc tgcgtc 196

<210> 729
 <211> 218
 <212> nucleic acid
 <213> Zea mays
 <400> 729

gattgactct gttctggatc ttgcacggTg cacgatcgTg tccgagctct ccaacaagga 60
 ttttgactcc tatgttctct ccgagtcaag cttgttcatc tatcctctga agattgtcat 120
 caagacctgt ggcactacca agctcctgct cacaattcca aggatectag agcttgctga 180
 agagctgtct atgcctcttg ctgctgtgaa gtactccc 218

<210> 730
 <211> 261
 <212> nucleic acid
 <213> Zea mays
 <400> 730

tggntgttaa tntnntcTnc acagccagta ccanaccaa ntangncng acnacncacc 60
 naccgntctct gtcttatggT gaccttgTca agagggtcct ccggtgcttt ggccccctga 120
 gttttccgTt gccgtgacca tcttcggcg gcgTggccat gccgggacat ggggaaagca 180
 cttggTgcag aggtctatga ctgcaacaac atggTggagc aggagctgcc tggagggggc 240
 tctcgtgta ccagagctna c 261

<210> 731
 <211> 306
 <212> nucleic acid
 <213> Zea mays
 <400> 731

atcaggggTt tcacacatag cttcgtcgat ttgaatttga tgtactaatg gagtctaagg 60
 gtggcaaaaa gtctagcagt agtcgtctat gatgtatgaa gctccccttg gctacagcat 120

tgaggacggtt cgacctgccg gagngtgaag aagttccagt ctgctgctta ctccaactgc 180
 gcgaagaagc catcctgata tcctttggct tcctcatcta gtagtttagg attctttctg 240
 acatttgatt ctgacaatct cctggctgtg ttctgataac gacaatcnca tctgtcttga 300
 tctcct 306

<210> 732
 <211> 295
 <212> nucleic acid
 <213> Zea mays
 <400> 732

ggacgttoga cctgccggan ggcgaagaag ttccagtctg ctgcttactc caactgcgca 60
 aagaagccat cctgatatcc cttttggctt cctcgttcta gtagtttagg atttcttttc 120
 tgacaottcg attcttacca atccccctgg cctgcngctt tcctgataat cgaccagttc 180
 cccagtcttg ctccctgcag tcctccctcc tccatctcca gcgttggtgt ctgactcacc 240
 tgcaccaatg gctgttcttt ctgctgctgg tgctccccgg gctnaagtat ngggt 295

<210> 733
 <211> 532
 <212> nucleic acid
 <213> Zea mays
 <400> 733

ggnnnnnnna actgtctatt cngcccgggc ccggtccaca aattnacggg gtcnacgcag 60
 cgteccgcnca cgcgctccgc ccacgccgtc cgcccacgcg tccggggcaa atgcttatgt 120
 gattggagat gcagcaagaa ccagggacag aagtggcaca tctactacgc cactgagacc 180
 cagagcaacc aatggtcaac cttgagatgt gcatgactgg tctggacacg aagaaacttc 240
 agtcttcttc aagactaatg ctgatggcaa cacaacatgt gccaaaggaaa tgacgagctc 300
 tctggttaanc tctgaaatta tcctgagat ggagatctgt gatcttgact tcgaaccctg 360
 cggtactcc catgaatgca atccatgggc tctgcgttct ccacgattca tgtgacgcct 420
 gaggatgggt tcaagctacg ccaagttatg aagttatggg gcntggatgc catggncctg 480
 ttctaaaggn gaaccttttc aaaaaggggc ccttnggggg cttccgcccc cc 532

<210> 734

<211> 269
 <212> nucleic acid
 <213> Zea mays

 <400> 734

 atccccctgg cctgctgctt tcttgacaat cgaccagttc cccagtcttg ctccctgcag 60
 tctccctcc tccatctcca gcgttggtgt ctgactcacc tgcaccaatg gctgttcttt 120
 ctgctgctgg tgctcccccg gcttcagcta tcgggtttga gggctatgag aagcgcttg 180
 agatcacatt ctctgaggca cctgtctttg tggaccccca tggcagcggg ttgcgtgccc 240
 tctccaggtc ccagattgac tctgttctg 269

<210> 735
 <211> 231
 <212> nucleic acid
 <213> Zea mays

 <400> 735

 gcttcgtoga tttgaatttg atgtactaat ggagtctaag ggtggcaaaa agtctagcag 60
 tagtcgttcc atgatgtatg aagctcccct tggctacagc attgaggacg ttcgacctgc 120
 cggagggtga agaagttcca gtctgctgct tactccaact gcgcgaagaa gccatcctga 180
 tatccctttt ggcttccgca tnnaggagtt agggtttcta cccnnccngn g 231

<210> 736
 <211> 220
 <212> nucleic acid
 <213> Zea mays

 <400> 736

 ogtttggtta ttctgtgaag cagccaagcc aggctattgt tatgaaaatt tgcgtctgt 60
 aagcatgtga acttccgatg ttgccacatg ctggatcagt ctgaataagt aagtatgcag 120
 ctctaggtgg tcagctgcgt ctaccacaat gagcatgaac gtatggagaa atatctgtga 180
 acccattttg tttatgaata agatttggtt ttttcgagtt 220

<210> 737
 <211> 442
 <212> nucleic acid
 <213> Zea mays

<400> 737

caaagaattc cctcccctag ccgccgncgc cgccgctcgt ctccctgaga ggtgccggga 60

ccgaggaact ggccgagtag taggtagaga atcgggtttt cttgtctcag cccgggggtct 120

gctgcgtctg gtgggtggtga aggggagaaa ttcgtgagat ctgttccgga tcaggcgtgc 180

gagctcggga atcaggggtt tcacacatag cttcgtcgat ttgaatttga tgtactaatg 240

gagtctaagg gtggcaaaaa gtctagcagt agtcgttcta tgatgtatga agctcccttg 300

gctacagcat tgaggacgtt cgacctgccg gaggcntnaa gaagttccag tctgctgctt 360

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ngattctttt ctgacctttg at 442

<210> 738

<211> 527

<212> nucleic acid

<213> Zea mays

<400> 738

gngngnggtg gtnanngnnn tcnannaana atctnctacc ggcccgggaat tcccgggtcg 60

acccacgcgt ccgaaagaat tccctcccct agccgccgcc gccgctcgtc tccctgagag 120

gtgccgggac cgaggatttg gncgagttct aggtagagaa tcggttttac ttgtctcagc 180

ccgggggtctg ctgcgtctgg tgggtggtgaa ggggagaaaat tcgtgagatc tgttccggat 240

caggcgtgcg agctcgggaa tcaggggttt cacacatagc ttcgtcgatt tgaatttgat 300

gtactaatgg agtctaaggg tggcaaaaag tctagcagta gtcgttctat gatgtatgaa 360

gctncccttg gctacagcat tgaagacgtt cgacctgccn gaagcctgaa aaagttcaat 420

ctgntgctta ctccaactgc gccaaagaaa ccattctgat atcccttttg gcttccctcat 480

tctaagtaag tttaaggatt cttttctgac actttgattc ttancaa 527

<210> 739

<211> 249

<212> nucleic acid

<213> Zea mays

<400> 739

ccttgggtgc ttcggcccct ccgaattttc tgtcgccgtg accatcttcg gcgggcccgg 60

ccaagctggg acatggggaa aggaacttgg tgcggaggct tatgactgca acaacatggt 120
cgagcaggag ctgcctggag gtgggatcct catctaccag agcttctgtg ctgctgaaga 180
cgccgttgct agctcgccca aatccgttct tcgctgcttt gatggcgaga atgcagcact 240
tttgcaag 249

<210> 740
<211> 514
<212> nucleic acid
<213> Zea mays
<400> 740

aagtgttnt ntngggggg gggnnntna atgttgacc ggtccggaat tcccgggtcg 60
accacgcgt ccggttgcca aagaattccc tcccctagcc gccgcgcgcg ctgctctccc 120
tgagaggtgc cgggaccgag gatttgccg agttctaggt agagaatcgg ttttacttgt 180
ctcagcccg ggtctgctgc gtctggtggt ggtgaagggg agaaattcgt gagatctgtt 240
ccggatcagg cgtgcgagct cgggaatcag gggtttcaca catagcttcg tcgatttgaa 300
tttgatgtac taatggagtc taagggtggc aaaaagtcta gcagtagtcg ttctatgatg 360
tatgaagctc cccttggtcta cagcattgag gacgttcgac ctgccggagg cgtgaagaag 420
ttccagtctg ctgcttactc caactgcgcg aagaaccatc ctgatatccc ttttggcttn 480
ctcattctag taanttaaga attcntttnt tgac 514

<210> 741
<211> 298
<212> nucleic acid
<213> Zea mays
<400> 741

gtagtttagg atttcttttc tgacacttcg attcttacca atccccctgg cctgctgctt 60
tcctgacaat cgaccagttc ccagtccttg ctccctgcag tctccctcc tccatctcca 120
gcgttggtgtt ctgactcacc tgcaccaatg gctgttcttt ctgctgctgg tgctcccccg 180
gcctcagcta tcgggtttga gggctatgag aagcgcttg agatcacatt ctctgaggca 240
cctgtctttg tggaccccat ggcagcggtt tgcgtgccct ctccaggtec cagattga 298

<210> 742

<211> 217
 <212> nucleic acid
 <213> Zea mays

<400> 742

ttttttttta actcgaaaaa aacaaatctt attcataaac aaaatgggtt cacagatatt 60
 tctccatagc ttcatgctca ttgtggtaga cgcagctgac cacctagagc tgcatactta 120
 cttattcaga ctgatccagc atgtggcaac atcggaagtt cacatgctta cagacgacaa 180
 attttcataa caatagcctg gcttggctgc ttcacag 217

<210> 743
 <211> 181
 <212> nucleic acid
 <213> Zea mays

<400> 743

tgcgattttg acttcgaacc ctgtggctac tccatgaatg cgatccatgg ctctgcattc 60
 tccacaatcc atgtgacgcc cgaggacggt ttcagctacg ccagttacga gggttatgggc 120
 ttggatgcca ctgctctgtc ttatggtgac cttgtcaaga gggtcctccg gtgctttggc 180
 c 181

<210> 744
 <211> 373
 <212> nucleic acid
 <213> Zea mays

<400> 744

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<210> 745

<211> 281
 <212> nucleic acid
 <213> Zea mays

 <400> 745

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 tgaatttnat gtactaatgg agtctaaggg tngcaaaaag tctagcagta gtcgtttctat 180
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 gaagttccag tctgctgctt actccaactg cgcgaagang c 281

<210> 746
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 <212> nucleic acid
 <213> Zea mays

 <400> 746

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<210> 747
 <211> 170
 <212> nucleic acid
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 <400> 747

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<210> 748
 <211> 441
 <212> nucleic acid
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 <400> 748

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 ggtggtggtg aangggagaa attcntgaga tctgtncogg atcaggcgng cgagctcggg 180
 aatcaggggt ttcacacata gcttcgtccg atttgaattt gatgtactan atggantcta 240
 anggtggcaa aaagtctanc attagtcgct ctatgatgta tgaanctccc ctnggctaca 300
 gcattgagga cgttcgacct gccggaggcg tgnanangtt ccagtctgat gcttactnca 360
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<210> 749
 <211> 195
 <212> nucleic acid
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<400> 749

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<210> 750
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 <212> nucleic acid
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<400> 750

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<400> 751

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cttgtggacc tcatgggcgt ggtttgctg nccctctnch agggcccaga ttgaactctg 180
tacnggatct tggcaggng caaaattgtg tctggagctc tccaacaagg atttcgactc 240
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<210> 752
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<212> nucleic acid
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<400> 752

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agcagtagtc gntccatgat gtatgaagct ccccttggt acagcatnng aanncgttcg 180
acctgccgga gnncaagaa gttccagtct gctgcttact ccaactgcgc aaagaagcca 240
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<210> 753
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<400> 753

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tgctgcgtag ggtggtgtg aaggggagaa gtttaagatc tgttccacag atcacgcgtg 240

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tggctacagc attgaggacg ttcgacctgc cggaggcgcc aagaagttcc aatctgctgc 420
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<210> 754
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<400> 754

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cccttttggc ttctcattc tagnagtta ggatttcttt totgacactt tgatgctgac 180
naatctctct ggctgggtgc ttctgataa tcgaccagtt cccagtcttg ctcttgcat 240
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<210> 755
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<212> nucleic acid
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<400> 755

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gctctaggtg gtcagctgcg tctaccacaa tgagcatg 158

<210> 756
<211> 165
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<400> 756

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tgctgatgct tccccggtct cagctatcgg gtttgagggc tatgagaagc gccttgagat 120

cacattctct gaggcacctg tctttgtnga ccttcattcn cntng 165

<210> 757
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 <212> nucleic acid
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<400> 757

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<210> 758
 <211> 221
 <212> nucleic acid
 <213> Zea mays

<400> 758

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<210> 759
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 <212> nucleic acid
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<400> 759

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 cgggaccgag gaactggccg agtactaggt agagaatcgg ttttcttgt ctcagccggg 180
 gtctgctgcg tctggtggtg gtgaagggga gaaattcgtg agatctgttc cggatcggcg 240
 tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc gatttgaatt tgatgactaa 300
 tggagtctaa ggggtggcaa aagtctagca gtagtcgttc taatgatgta tgaactcccc 360
 ttggctacag cattgangac gttcgacctg ccggaggcgt naagaanttc cagcctgctg 420

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<210> 760
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tcgttttatg atgtatgaag ctccccttgg gctacagcat tgaggangtt cgacctgccg 180
gacgtgaaga agttccagtc tgntgnttac tccaaatg 218

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ttcggcgggc gtggccatgc cgggacatgg ggaaaggcac ttggtgcaga ggtctatgac 120
tgcaacaaca atggtggaat aggancttcc tggaaggcgg ggntcctcgt gtaacaaaaa 180
acttctgtgc tgctnaagac ncngttgcta acctcgcccc aaattttt 228

<210> 764

<211> 147

<212> nucleic acid

<213> Zea mays

<400> 764

agcttgatga gtaagacggg cttctggtgt cgatttgctt ctgagttgtt tattttatat 60
cgtcgcaatt tcgtggttgt cgtttggtta ttctgtgaag cagccaagcc aggctattgt 120
tatgaaaatt tgctgtctgt aagcatg 147

<210> 765

<211> 187

<212> nucleic acid

<213> Zea mays

<400> 765

cgtgaccatc ttcgccggg cgtggccatg ccgggcacat ggggaaaggc actaggtgca 60
gaggtctatg actgcaacaa catgtggagc aggagctgcc tggaggcggg ctctcgtat 120
accagagctt ctgcgctgct gaagacgctg ttgctacctt cgcccaaate tgttttccac 180
tgctttg 187

<210> 766

<211> 154

<212> nucleic acid

<213> Zea mays

<400> 766

gcttctgat aatcgaccag ttccccagtc ttgctccttg cactcctccc tctccatct 60
ccagcattgt gttctgattc acctgctcca atggctgttc tttctgctgc tgatgcttcc 120
ccggtctcag ctatcgggtt tgaggcatg agaa 154

<210> 767
 <211> 162
 <212> nucleic acid
 <213> Zea mays

 <400> 767

 tctggatctt gcacggtgca caattgtgtc cgagctctcc aacaaggatt tcgactcata 60
 tgtacctctc tgagtcaagc ttgtttatct atccctctga agattgtnat caagaacctgt 120
 ggcactacca agctcctgct caccattcca agaatccttg ag 162

<210> 768
 <211> 320
 <212> nucleic acid
 <213> Zea mays

 <400> 768

 ccctccccta gccgccgccg ccgctcgtct ccctgagagg tgccgggacc gaggaactgg 60
 ccgagttcta ggtagagaat cggttttgct tgtctcagca cggggtctgc tgcgtctggt 120
 ggtggtgaag gggagaaaatt cgtgagatct gttccggatc aggcgtgcga gctcgggaat 180
 cagggggtttc acacatagct tcgtcgattt gaatttgatg tactaatgga gtctaagggt 240
 ggcaaaaagt ctagcagtag tcgttccatg atgtatgaag ctccccttgg ctacagcatt 300
 gaggacgttc gacctgccgg 320

<210> 769
 <211> 295
 <212> nucleic acid
 <213> Zea mays

 <400> 769

 ctcgtctccc tgagangtgc cgggaccgag gaactggccg agttctaggt agagaatcgg 60
 ttttgcttgt ctcagcannn ggtctgctgc gtctggtggt ggtgaagggg agaaattcgt 120
 gagatctggt ccggatcagg cgtgcgagct cgggaatcag gggtttcaca catagcttcg 180
 tcgatttgaa tttgatgtac taatggagtc taagggtggc aaaaagtcta gcagtagtcg 240
 ttccatgatg tatgaagctc cccttggtc cagcattgag gacgttcgac ctgcc 295

<210> 770

<211> 329
 <212> nucleic acid
 <213> Zea mays

 <400> 770

 ccatcttcgg cgggcgtggc catgncgggg acatggggaa aggacttggt gnagaggtct 60
 atgactgcaa caacatgggt gagcagagct gcctggagcg ggctctctgt accagagttc 120
 ngtgctgctg aagacgtgtg ntactcgcca atctgttncc actgcttgac ggcgagacgt 180
 ggaagtgtcc tcctcctatg aagaaggata caagctggct aatcttctct gtgggaggag 240
 gagcggatnc atggaggaga aggcgggatg cttgattgag tagacgggtt ctggggtcga 300
 ttnttctgag tgttattnn cgcgcaatt 329

<210> 771
 <211> 511
 <212> nucleic acid
 <213> Zea mays

 <400> 771

 agnaggtggt ttaattagnn gaagggnggn aatgaaaga ctntccggct ngaattcccg 60
 ggtcgaccca cgcgtccgat tnantcccat agccgacnac ggcgntcgtc tgctgagag 120
 gtgccggggac cgaggnactg gncgagttct agggagagaa tcggttttgc ttgtctnagc 180
 acggggtctg ntgcgtctgg tgggtggtgaa ggggagaaat tcgtgagatc tgttccggat 240
 caggcgtgcg agctcgggaa tcaggggttt cacacatagn ttcgtngatt tgaatttgat 300
 gtactaatgg agtctaaggg tggcaaaaag tctagcangg agtcgttcca tgatgtatga 360
 agtccccctt ggntacagga ttgaagacgt tagactncct ganggcatga aaaaagttca 420
 agctggnggt tnnttcaanc nccncnanaa accgttctn aattcccttt nggttnctta 480
 atctnaaann ttaaganttn ctttcttaaa a 511

<210> 772
 <211> 154
 <212> nucleic acid
 <213> Zea mays

 <400> 772

 taagcatgtg aacttccgac gttgccacat gctggatcag tctgaataag taagtatgca 60

gctctaggtg gtcagctgcg tctaccacaa tgagcatgta cgtatggaga aatatctgtg 120
 aaoccatttt gtttatgaat aagatttggtt tttt 154

<210> 773
 <211> 160
 <212> nucleic acid
 <213> Zea mays
 <400> 773

taagcatgtg aacttccgat gttgccacat gctggatcag tctgaataag taagtatgca 60
 gctctaggtg gtcagctgcg tctaccacaa tgagcatgac gtatggagaa atatctgtga 120
 aoccattttg tttatgaata agatttggtt ttttcgagtt 160

<210> 774
 <211> 126
 <212> nucleic acid
 <213> Zea mays
 <400> 774

cctccctcca tctccagcat tgtgttctga ttcacctgct ccaatggctg ttctttctgc 60
 tgctgatgct tccccggtct cagctatcgg gtttgnnggc tatgngaagc gccttgagat 120
 cacatt 126

<210> 775
 <211> 391
 <212> nucleic acid
 <213> Zea mays
 <400> 775

aatctctacn gctccagggc ccggnccagg aantacgng tcgacccatc cacacaggca 60
 nccggttgcc aaagaattcc ctcccctagc cgccgccgcc gctcgtctcc ctgagaggtg 120
 ccgggaccga ggaactggcc gagttctagg tagagaatcg gttttgcttg tctcagcacg 180
 gggctctgctg cgtctgggtg tgggtgaagg gagaaattcg tgagatctgt tccggatcag 240
 gcgtagcgagc tcgggaatca ggggtttcac acatagcttc gtcgatttga atttgatgta 300
 ctaatggagt ctaagggtgg caaaaagtct agcagtagtc gttocatgat gtatgaagct 360
 ccccttggtc acagcattga ggacgttcga c 391

<210> 776
 <211> 493
 <212> nucleic acid
 <213> Zea mays

 <400> 776

 gtgagngttt ntttngnnna nggnggngng atgntatctt cgtccggaat tcccgggtcg 60
 acccacgent ccgctnccct agccgccncc gcccgntcgt ctccctgaga ggtgccngga 120
 ccgangaact ggccgngttc taggtagaga atnggttttg cttgtctcan cacggggtct 180
 gctgcgtctg gtgggtggtga aggggagaaa ttcgtgagat ctgttccgga tcaggcgtgc 240
 gagctcggga atcaggggtt tcacacatag ctctcgtcgt ttgaatttga tgtactaatg 300
 gagtctaagg gtggcaaaaa gtctagcagt antcnttcca tgatgtatga agctncccnt 360
 ggctacagca ttgaggacgt cnanctnccc ggaagcntta aanaattnca gnctgtgntt 420
 antnnaactg nnaanaaac nattctgana tnnctttttg gntttnttan tnnaanantt 480
 nanganttnt ttt 493

<210> 777
 <211> 193
 <212> nucleic acid
 <213> Zea mays

 <400> 777

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 agctncnggg ttggtcagnt gcgtctagca naatgagcat gtacgtatgg agaaatntnn 120
 tgtgaacca ttttgtttat gaataagatt tgtttttttt cgagttaaaa naaaaaaaaa 180
 agggcgccn cng 193

<210> 778
 <211> 136
 <212> nucleic acid
 <213> Zea mays

 <400> 778

 gtgacgcccg aggacggttt cagctatgcc agttatgagg ttatgggctt ggatgccact 60
 gctctntcct tatggtgacc ttgtcaagag ggtccttcgg tgctttggcc cctcggagtt 120

ttccgttgcc gtganc

136

<210> 779
<211> 117
<212> nucleic acid
<213> Zea mays

<400> 779

ggctgttctt tctgctgctg atgcttcccc ggtctcagct atcggggttg agggctatga 60

gaagcgcctt gagatcacat tctctgaggc acctgtcttt gtggaccctc atgggcg 117

<210> 780
<211> 253
<212> nucleic acid
<213> Zea mays

<400> 780

tcccctagcn gccgcgcgcg ccgatcgtct cccctgagag gtgccgggac cgangaactt 60

ggccgagtag taggtagaga aattcgtgag atctgttccn gatcangcgt gcgagctcgg 120

gaatcagggg tttcacacat agctcgtcga tntgaatttg atgtactaat ggagtctaag 180

ggtggcaaaa agtctagcag tagtcgttct atgatgtatg aagctcccct tggctacagc 240

attgatgacg ttc 253

<210> 781
<211> 273
<212> nucleic acid
<213> Zea mays

<400> 781

tcgtctccct gagaggtgcc gggaccgagg aactggccga gtactaggta gagaatcgg 60

ttttcttgct tcagcccggg gtctgctgctg tctggtggtg gtgaagggga gaaattcgtg 120

agatctgttc cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt 180

cgatttgaat ttgatgtact aatggagtct aagggtggca aaaagtctag cagtagtcgt 240

tctatgatgt atgaagctcc ccttggttac agc 273

<210> 782
<211> 110
<212> nucleic acid

<213> Zea mays

<400> 782

gtggagagtg ctctctctcc tatgaagaag gactacaagc tggctaattct tctctgctgg 60

gaggaggaag cggatgccat ggaggagaag gcgggagtgcc ttgatgagta 110

<210> 783

<211> 272

<212> nucleic acid

<213> Zea mays

<400> 783

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ttttcttgtc tcagcccggg gtctgctgcg tctggtggtg gtgaagggga gaaattcgtg 120

agatctgttc cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt 180

cgatttgaat ttgatgtact aatggagtct aagggtgcca aaaagtctag cagtagtcgt 240

tctatgatgt atgaagctcc ccttggtctac ag 272

<210> 784

<211> 491

<212> nucleic acid

<213> Zea mays

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tctaggtaga gaatcggttt tgcttgtctc agcacggggg ctgctgcgtc tgggtggtgg 180

gaaggggaga aattcgtgag atctgttccg gatcaggcgt gcgagctcgg gaatcagggg 240

tttcacacat agcttcgtcg atttgaattt gatgtactaa tggagtctaa ggggtggcaa 300

aagtctaaca agtagtcgtt ccatgatgta tgaagctccc cttggctaca gcatttgaag 360

gacgttcgaa cctgccggaa gcnttnaaga agtttncaagt ctgctgntta cttccaactt 420

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ttcctttctt g 491

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 <212> nucleic acid
 <213> Zea mays
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 ttttcacagc ctgggggtctg ctgcgtaggg tgggtgtgaa ggggagaagt ttaagatctg 120
 ttccacagat cagcggtgcg cgctcgcgaa tcgggggttc cacacatagc ttcgtcgttt 180
 tgaatttgat gtactaatgg agtctaangg tggcaagaag tctagcagta gtcgttccat 240
 gatgtatgaa gctccccttg gctacagcat tga 273

<210> 786
 <211> 310
 <212> nucleic acid
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 ctgctgcgtc tggnggtggg gaaggggaga aattcgtgag atctgttccg gatcaggcgt 180
 gcgagctcgg gaatcagggg tttcacacat agcttcgtcg atttgaattt gatgtactaa 240
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 ttggctacag 310

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 <212> nucleic acid
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 ctgctgctgg tgggtgtgaa ggggagaaat tcgtgagatc tggtccggat caggcgtgcg 180
 agctcgggaa tcaggggttt cacacatagc ttcgtcgatt tgaatttgat gtactaatgg 240
 agtctaaggg tggcaaaaag tctagcagta gtcgttccat gatgtatgaa gctccccttg 300

gctacag 307

<210> 788
<211> 107
<212> nucleic acid
<213> Zea mays

<400> 788

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cctgggtgcac agccagcccc ccacaggagc ttctctgagg aagttgc 107

<210> 789
<211> 265
<212> nucleic acid
<213> Zea mays

<400> 789

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ttttctttgtc tcagcccggtg gtctgctgctg tctgggtggtg gtgaagggga gaaattcgtg 120
agatctgttc cggatcaggg gtgagagctc gggaatcagg ggtttcacac atagcttcgt 180
cgatttgaat ttgatgtact aatggagtct aagggtggca aaaagtctag cagtagtcgt 240
tctatgatgt atgaagctcc ccttg 265

<210> 790
<211> 304
<212> nucleic acid
<213> Zea mays

<400> 790

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gggtttcaca catagcttcg tcgatttgaa ttgatgtac taatggagtc taaagggtgg 240
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<211> 149
 <212> nucleic acid
 <213> Zea mays

<400> 791

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 <212> nucleic acid
 <213> Zea mays

<400> 792

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 ttccacagat caagcgtgcg cgctcgcgaa tcgggggttc cacacatagc ttcgtcgttt 180
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 <212> nucleic acid
 <213> Zea mays

<400> 793

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 <212> nucleic acid
 <213> Zea mays

<400> 794

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aatcctgtg acgcc 135

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<211> 258

<212> nucleic acid

<213> Zea mays

<400> 795

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<212> nucleic acid

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<212> nucleic acid

<213> Zea mays

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 tctatgatgt atga 254

<210> 800
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 <212> nucleic acid
 <213> Zea mays
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tctatgatgt atgaa 255

<210> 801
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<212> nucleic acid
<213> Zea mays

<400> 801

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<210> 802
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<212> nucleic acid
<213> Zea mays

<400> 802

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tctatgatgt atgaa 255

<210> 803
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 803

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ctgctgcgtc tgggtggtgt gaaggggaga aattcgtgag atctgttccg gatcaggcgt 180

gcgagctcgg gaatcagggg ttccacacat agctccgtcg atttgaattt gatgtactaa 240

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 <211> 295
 <212> nucleic acid
 <213> Zea mays

<400> 804

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cgatttgaat ttgatgtact aatggagtct aagggtgcaa aaagtctagc agtagtcgtn 240

tatgatgtat aaagcncccc ttggctanag cattgaggac gttngacctg ccggg 295

<210> 805
 <211> 255
 <212> nucleic acid
 <213> Zea mays

<400> 805

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ttccacagat cagcgtgctg cgctcgcgaa tcgggggttc cacacatagc ttcgtcgttt 180

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gatgtatgaa gctcc 255

<210> 806
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 <212> nucleic acid
 <213> Zea mays

<400> 806

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aggggagaaa ttccngagat ctgttccgga tcaggcgtgc nantcggna atcagggggtt 180

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gtctagcagt agncgttcta tgatgtatga agctcccc 278

<210> 807
<211> 127
<212> nucleic acid
<213> Zea mays

<400> 807

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tgtntga 127

<210> 808
<211> 251
<212> nucleic acid
<213> Zea mays

<400> 808

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tctatgatgt a 251

<210> 809
<211> 250
<212> nucleic acid
<213> Zea mays

<400> 809

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tctatgatgt 250

<210> 810

<211> 250
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 <213> Zea mays
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 tctatgatgt 250

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<210> 811
 <211> 260
 <212> nucleic acid
 <213> Zea mays
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 ttccacagat cagcgtgctg cgcttcgcca atcggggggtt ccacacatag ctctgctggt 180
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<210> 812
 <211> 257
 <212> nucleic acid
 <213> Zea mays
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<210> 813

<211> 155
 <212> nucleic acid
 <213> Zea mays

<400> 813

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 ggotgttctt tctgctgctg gtgtccccc gntca 155

<210> 814
 <211> 276
 <212> nucleic acid
 <213> Zea mays

<400> 814

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 aagggtggca aaaagtctag cagtagtctg tccatg 276

<210> 815
 <211> 97
 <212> nucleic acid
 <213> Zea mays

<400> 815

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 <211> 262
 <212> nucleic acid
 <213> Zea mays

<400> 816

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 aaagtctagc agtagtcgtt cc 262

<210> 817
 <211> 281
 <212> nucleic acid
 <213> Zea mays
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 <211> 250
 <212> nucleic acid
 <213> Zea mays
 <400> 818

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<210> 819
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gagatctgtt ccgcatcagg cgtgcgagct cgggaatcag gggtttcaca catagcttcg 180
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 tctatgatgt atg 253

<210> 820
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 <212> nucleic acid
 <213> Zea mays
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 <400> 822

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cgatttgaat ttgatgtact aatggagtct aagggtnngn aaaagtctag cagtagtcgt 240
ctatgatgta tna 253

<210> 823
<211> 269
<212> nucleic acid
<213> Zea mays

<400> 823

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tctaagggtg gcaaaaagtc tagcagtag 269

<210> 824
<211> 296
<212> nucleic acid
<213> Zea mays

<400> 824

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gtactaatgg agtctaaggg tggcaaaaag tctagcagta gtcttccatg atgtat 296

<210> 825
<211> 296
<212> nucleic acid
<213> Zea mays

<400> 825

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<210> 826
 <211> 267
 <212> nucleic acid
 <213> Zea mays

<400> 826

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 <212> nucleic acid
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<400> 827

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 <211> 295
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<213> Zea mays

<400> 831

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<210> 832

<211> 227

<212> nucleic acid

<213> Zea mays

<400> 832

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gatctgttcc ggatcaggcg tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc 180
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<210> 833

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 833

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<210> 834

<211> 222

<212> nucleic acid

<213> Zea mays

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 <211> 261
 <212> nucleic acid
 <213> Zea mays

<400> 835

aaagaattcc ctcccctagc cgcgcngcc gctcgtctcc ctgagaggtg ccgggaccga 60
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 cgtctgggtg tggtgaagg gagaaattcg tgagatctgt tccggatcag gcgtgcgagc 180
 tcgggaatca ggggtttcac acatagcttc gtcgatttga atttgatgta ctaatggagt 240
 ctaagggtgg caaaaatcta c 261

<210> 836
 <211> 97
 <212> nucleic acid
 <213> Zea mays

<400> 836

tcagctgcgt ctaccacaat gagcatgaac gtatgngac atatctgtga acccattttg 60
 tttatgaata agatttggtt ttntcgagtn aaaaaaa 97

<210> 837
 <211> 229
 <212> nucleic acid
 <213> Zea mays

<400> 837

gtctccctga gaggtgccgg gaccgaggaa ctggccgagt tctaggtaga gaatcggttt 60
 tgcttgtctc agcacggggt ctgctgcgtc tgggtgggtg gaaggggaga aattcgtgag 120
 atctgttccg gatcaggcgt gcgagctcgg gaatcagggg tttcacacat agctcgtcga 180
 tttgaatttg atgtactaat ggagtctaag ggtggcaaaa agtctagca 229

<210> 838
 <211> 254
 <212> nucleic acid
 <213> Zea mays

 <400> 838

 ganttcctc ccctagccgc cgcgcgcgct acgtctccct gagagggtgcc gggancgagg 60
 aactggccga gttctaggta gagaatcggg tttgcttgct tcagcacggg gtctgctgcg 120
 tctgggtggtg gtgaagggga gaaattcgtg agatctgttc cggatcaggc gtgcgagctc 180
 gggaatcagg ggtttcacac atagcttcgt cgatttgaat ttnatgtact aatggagtct 240
 aagggtggca aana 254

<210> 839
 <211> 493
 <212> nucleic acid
 <213> Zea mays

 <400> 839

 caaagaattc cctcccctag ccgcgcgncgn cgctcgtctc cctgaagaag gtgcccggga 60
 ccgagggatt tggcccagat tctaggtaag aagaaatcgg gttttacttg tctcaagccc 120
 ggggggtctgc ttgcgtctgg tgggtgggtga aaggggaaga aattcgtgaa gatctgttcc 180
 ggatcaggcg tgcgaanctc gggaatcagg gggtttcaca cataagcttc gtcgatttga 240
 atttgatgta ctaaattgga agtctaaagg gtggcaaaaa gtctagcagt aagtccgttc 300
 tatgatgtat gaaacttccc tttggctaca gcatttgaag gacgttcgac ctggccggan 360
 gccttgaaga angttccagt ctgnttgntt acttcaactt gngccaaaaa accattctgg 420
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 aatcttganc caa 493

<210> 840
 <211> 296
 <212> nucleic acid
 <213> Zea mays

 <400> 840

 angnacgcgt acgttagctc gnaattcngc tcgagcggct cgagcnaaga attccntccc 60

ctagccgccg ccgncgccgc tcgtctccct gagaggtgcc ggnaccgagg aactggccga 120
 gtactaggta gagaatcgnt tttcttctgc tcagcccggg gtctgtctgc tctggtggtg 180
 gtgaanggga gnaattcgtn agatctgttc cggatcaggc gtgcgagctc gggaatcagg 240
 ggtttcacac atagcttcgt cgatttgaat ttgatgtact aatggagtct aagggg 296

<210> 841
 <211> 292
 <212> nucleic acid
 <213> Zea mays
 <400> 841

tngccnnana nttccnnccc ctagccgccn ccgcccgcgc tngtntccct gagaggtanc 60
 cgggaccgat gaactngccg agtactaggt agagaatcgg tttntcttnt ctcagcccgg 120
 ggtctgctnc ntctggtggt ggtgaagggg agaaattcgt ganatctgtt ccggancatg 180
 cgtgngagct cnggaatcag gggtttcaca catagcttcg tcgatttgaa tntgatgtat 240
 aatggatcta agggtnngcaa naantctagc agtagtcgtn cnagntgnat ga 292

<210> 842
 <211> 254
 <212> nucleic acid
 <213> Zea mays
 <400> 842

cgccgccgct catctcgccg agaggtgacg ggaccgagga actcgccgag ttctaagata 60
 gagaatcgct tttcacagcc tggggtctgc tgcgtagggg ggtgttgaag gggagaagtt 120
 taagatctgt tccacagatc acgcgtgcgc gctcgcgaa cgggggttcc acacatagct 180
 tcgtcgtttt gaattgatgt actaatggag tctaaagggtg gcaagaagtc tagcatatcg 240
 ttccatgatg tatg 254

<210> 843
 <211> 255
 <212> nucleic acid
 <213> Zea mays
 <400> 843

gccaaagaat tccctccctt agccgccgcc gccgctcgtc tccctgagag gtgccgggac 60

cgaggaactg gccgagttct aggtagagaa tcggttttgc ttgtctcagc acggggtctg 120
 ctgcgtctgg tgggtggtgaa ggggagaaat tcgtgagatc tgttccggat caggcgtgcg 180
 agctcgggaa tcaggggttt cacacatagc ttcgctgatt tgaattgatg tactaatgga 240
 gtctaagggt ggcaa 255

<210> 844
 <211> 214
 <212> nucleic acid
 <213> Zea mays
 <400> 844

tcctctcgcc gagaggtgac gggaccgagg aactcgccga gttctaagat agagaatcgc 60
 ttttcacagc ctggggtctg ctgcgtaggg tgggtgtgaa ggggagaagt ttaagatctg 120
 ttccacagat cagcgtgcg cgctnccga atcggggggt ccacacatag cttcgtcgtt 180
 tgaatttgat gtactaatgg agtctaaggg tggc 214

<210> 845
 <211> 798
 <212> nucleic acid
 <213> Zea mays
 <400> 845

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 gggggatcaa cattcttctt gagggcncct tggcttttg tgggaccct catgggcgtg 120
 ggttttgncg tgccctctt ccaaggcccc agattgactc tgttctggga tctttgcacg 180
 ggngcacaaa ttgtgtcccg agctcttcca acaagggtt tccgactcat atgtcctttc 240
 tgaggcaaag cttgnntatc tctctctga agattgtcat caaagacctg tggcactacc 300
 aagctcctgc tcaccattcc aaagaatcct tgaacttgct tgaaaaactt gtctatgcc 360
 ctttgctgct tgtgaaaata ctcccgtagg acggtcatct tttctggng cacagncagg 420
 cccccacaa ggaacttctt ttgaaggaag ttgntgnact ttaaaccgct tcttttggcc 480
 ggctggaaa tctggggggg naatgcntn tgtggaattg gggaatncca ncnaggaacc 540
 tggacaanaa atnggnaccg ntttttaacn ccccttgatt ncccnagca acccaatggg 600
 tnaacctna aaaggccnt gactgggctg ggcnnnnnnn nnnnnnnnnn nnnnnnnnaa 660

naataatgtt gangggaaca caacttgggc caagggaaat ganaaacttt ttgnnttttg 720
 aaaatatncc ccgaanggaa aacngnantt tantttaacc cgggggttttc aggaaggggc 780
 ccagggtttg ttttcnaa 798

<210> 846
 <211> 268
 <212> nucleic acid
 <213> Zea mays
 <400> 846

tcgtctccct gagaggtgcc gggaccgagg aactggccga gtactaggta gagaatcggg 60
 ttttcttgtc tcagcccngg gtctgtctgog tctgggtggtg gtgaanggga gaaattcgtg 120
 aganctgtac cactcaggcg tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc 180
 gatttgaatt tgatgtacna tggagtctaa ggggtggcaaa aatctagnon tatcgttcta 240
 ngattatnaa gccccctngg taaannat 268

<210> 847
 <211> 276
 <212> nucleic acid
 <213> Zea mays
 <400> 847

gcngccgccc cncctcgtct ccctgagagg tgccgggacc gaggaactgg ccgagtacta 60
 ggtagagaat cggtttttct tgtctcagcc cgggggtctgc tgcgtctggt ggtggtgaag 120
 gggagaaatt cgtgagatct gttccggatc angcgtgcga gtcgaggat caggggtttc 180
 acacatagct cgtcgatttg aatttgatgt acnaatggat cnaaggggtg caaaaagtct 240
 agcagnagtc gtctatgatg tatgaagctc cccttg 276

<210> 848
 <211> 185
 <212> nucleic acid
 <213> Zea mays
 <400> 848

gcaagctacc tgtgacaaaa acgattccaa gaaaccactt cttgtgcttc ctccactggc 60
 tgtctgtgca tcttgctcta aaacaatcgc agtcctgcaa gttgttgctg ttgctcctac 120

ccctgcctct gcaattgggt ttgagggata tgagaagcgc ctcgagatca gcttctatga 180
ggcac 185

<210> 849
<211> 519
<212> nucleic acid
<213> Zea mays
<400> 849

gnaagggatc tttnaaggng ggggggggga aaanganaac tncgtaccgg tccggaattc 60
ccgggtcgac ccacgcgtcc gagagottcg actcatatgt tctgtccgag tccagcctct 120
ttgtgtatcc gggacagggg tgtttctgaag acctgtggga cgacgaagct gctgctcgcc 180
attcctcgca tccttgagct tgotgcaggg ctgtcactgc ctcttctttc agcgaagtac 240
tctcgtggga ctttcatctt ccccggcgcg cagcccgcg caccgcgag cttctcggag 300
gaggatatctg tgctgaacgc tttcttttggg aacctcaagt ccggtggcaa tgcctacctg 360
atcgggtgaac cgttcaagcc caacaagaag tggcatgtct actacgccac agaggagcct 420
gagcgtccta tgggtgacgt tgagatgtgc atgacagagc ttgacgttaa gaaagctgct 480
gtgttcttca agaacttcac tgggtggcaac ttnacntna 519

<210> 850
<211> 324
<212> nucleic acid
<213> Zea mays
<400> 850

atgagaagcg cctcgagatc acgttctctg actcgctgt ctttgaggac ctttgtggtc 60
gtggcctgcg cgcctctctc cgtgagcaga tcgactcgtt tctggatctc gcacggtgca 120
ccatagtgtc acagctctcc aacgagagct tcgactcata tgttctgtcc gagtccagcc 180
tctttgtgta tccccacaag gttgttctga agacctgtgg gacgacgaag ctgctgctcg 240
ccattcctcg catccttgag cttgctgcag ggctgtcact gcctcttctt tcagcgaagt 300
actctcgtgg gaccttcac ttcc 324

<210> 851
<211> 260
<212> nucleic acid

<213> Zea mays

<400> 851

gaagcgccctc gagatcacgt tctctgacgc gcctgtcttt gaggaccctt gtggtcgtgg 60
cctgcgcgcc ctctcccgtg agcagatcga ctcgtttctg gatctcgac ggtgcacccat 120
agtgtcacag ctctccaacg agagcttcga tcatatgtt ctgtccgagt ccagcctctt 180
tgtgtatccc cacaaggttg ttctgaagac ctgtgggacg acgaagctgc tgctcgccat 240
tcctcgcatc cttgagcttg 260

<210> 852

<211> 272

<212> nucleic acid

<213> Zea mays

<400> 852

agcgccctcga gatcacgttc tctgacgcgc ctgtctttga ggacccttgt ggtcgtggcc 60
tgcgcgccct actcccgtga gcagatcgac tcgtttctgg atctcgacg gtgcaccata 120
gtgtcacagc tctccaacga gagcttcgac tcatatgttc tgtccgagtc cagcctcttt 180
gtgtatccca caaggttgtt ctgaagacct gtgggacgac gaagctgctg ctcgccattc 240
ctcgcatcct tgagcttgct gcaggctgtc at 272

<210> 853

<211> 239

<212> nucleic acid

<213> Zea mays

<400> 853

gcgcctcgag atcacgttct ctgacgcgcc tgtctttgag gacccttgtg gtcgtggcct 60
gogcgccctc tcccgtgagc agatcgactc gtttctggat ctgcgacggg gcaccatagt 120
gtcacagctc tccaacgaga gcttcgactc atatgttctg tccgagtcca gcctctttgt 180
gtatccccac aaggttgttc tgaagacctg tgggacgacg aagctgctgc tcgccattc 239

<210> 854

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 854

cctgcgcgcg ctnctncgtg agcagatcga gtcgtttctg gatctcgcac ggtgcaccat 60
agtgtctcag ctctnccaan gagagctnag actcatatgt tctgtncgag tccagcctct 120
ttgtgtatcc ccacaaggnt gttctgaaga cctgtggnac gacgaagctg ctgctcgca 180
ttcctcgcac ccttgagctt gctgcagggc tgtcantgcc tcttctttca gogaagtact 240
ctcgtgggac cttcatcttc cccggcg 267

<210> 855

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 855

attggctttg agggctatga gangcgcctc gagatcacgt tctctgacgc gcctgtcttt 60
gaggaccctt gtggctcgtg cctgcgcgcg ctctcccgtg agcagatcga ctggtttctg 120
gatctcgcac ggtgcaccat agtgtcacag ctctccaacg agagcttcga ctcatatgtt 180
cagtccgagt ccagcctctt gtgtatcccc acaaggttgt tncgaagact gtgggacgac 240
gaagctgctg ctgccattc ctgcacatnt gagntgctgc aggtgtcat gcctc 295

<210> 856

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 856

caccatagtg tcacanctct ccaacgagag cttcgactca tatgtttctgt ccgaatccag 60
cctctttgtg tatccccaca aggttggttct caagacctgt gggacnacga agctgctgct 120
cgccattcct cgcaccttg agcttgctgc agggctgtca ctgcctcttc tttcagcgaa 180
gtactctent gggaccttca tcttccccgg cgcgcagccg cgccacaccg cagcttctcg 240
gaggaggtat ctgtgctgaa cgcttt 266

<210> 857

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 857

cagcctcttt gtgtatcccc acaaggttgt tctgaagacc tgtgggacga cgaagctgct 60
gctcgccatt cctcgcatcc ttgagcttgc tgcagggctg tcaactgcctc ttctttcagc 120
gaagtactct cgtgggacct tcatcttccc cggcgcgcacn ccgcgccaca ccgcagcttc 180
tcggaggagg tatctgtgct gaacgctttc tttggtaacc tcaagtccgg tggcaatgcc 240
tacctgatcg gtgaaccgtt caagcccaac aagaagtggc atgtctacta cgccacagag 300
gagcctgagc gtccta 316

<210> 858

<211> 219

<212> nucleic acid

<213> Zea mays

<400> 858

ggtatcttaa acttagaact cttcagcgtg catcatatac tcattacttc tatgaagcaa 60
ctgttggtgc tggcctccct atcattagca ctctgcgtgg tctcctagag acaggtgaca 120
agatattgcy cattgaaggg attttcagtgc gaacgttaag ttatatTTTT aacaacttcg 180
aaggagcaag aaccttttagt gatgttggtg ctgaagcaa 219

<210> 859

<211> 323

<212> nucleic acid

<213> Zea mays

<400> 859

cagcactcca ggagtcagtgc caatactttt tgatgcatta gcaaaggcaa acattaatgt 60
tcgagcaata gctcaagggt gcagtgaata caatattact gttgtattga agcaacaaga 120
ttgtgtaagg gctctgagag ctgctcactc aaggttcttc ctctccaaaa ccacacttgc 180
tgttgggatac attggacctg gtttaattgg tggagcactc cttaaccagt tgaagaatca 240
gactgctggt ctcaaggaaa atatgaacat tgatctgcgt gtcattggaa taactggctc 300
aagtacaatg cttttgagcg aca 323

<210> 860

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 860

ctttctgata ttcctgtgaa gagccttgtg ccagaaacac ttgcagcatg ctgctcagca 60
gatgaattca tgcagaagct accatccttt gacgaggact gggccaggca acgcagttag 120
tgctgatgcn gcagacgaag tcctgcgtta cgttggagtg gtggatacgg tgaacaagag 180
gggacagggtg gagctacgta ggtacaagag ggaccaccgg ttgcgcant ctccggctcc 240
gacaacatca tcgcattcac cacctcgagg tacaaggagc agccgctaata cgtgcggngc 300
ccaggcgcgg 310

<210> 861

<211> 242

<212> nucleic acid

<213> Zea mays

<400> 861

tgaacactct gtatgctttg ctgttccaga aaaggaagtt gcattggttt ctgcagcctt 60
gcatgctagg ttctgtgaag cattggcagc agggagggtta tctaagggtg aagtcattcn 120
taattgtagc atccttgcta ctgtcggcct gaggatggca agtacacctg gagtcagcgc 180
gacccttttt gatgcactag ctaaggcaaa tnttaatgta agggcgatan tcaagggtgc 240
ag 242

<210> 862

<211> 303

<212> nucleic acid

<213> Zea mays

<400> 862

tgaaaaacga cttgaaaaat ggttttctcg atgtccggct gagacaataa ttgctactgg 60
gtttattgcg agtacacctg aaaacatccc aacaacctta aaaagagatg gaagtgattt 120
ctctgcagcc ataattggtt ccttggttaa ggctcgccag gtgaccatct ggactgatgt 180
tgatggggta tttagtgcag atcccagaaa agttagttag gctgtaatct taagcacatt 240
gtcgtatcag gaggcctggg aaatgtcata ttttggggca aatgtcttgc atccacgtac 300
tat 303

<210> 863
 <211> 294
 <212> nucleic acid
 <213> Zea mays

 <400> 863

 gtgaagtcag catatctttg aacttgatc catcgaaact atggagttgt gaattggttc 60
 agcacaaaaa tgaacttgat gatgtcattg aagagcttga gaagatagca gttgttcac 120
 tactgcagaa caggtcaata atatctttaa tttgaaatgt gcagaggtcg tctttgattc 180
 ttgaaaaggc tttcaatgtt ctacgaagaa acggtgtcaa tgtccagatg atatcgcaag 240
 gggcgtccaa ggtgaacatt tccttggtgg tccatgacag tgaggcaaaa cagt 294

<210> 864
 <211> 224
 <212> nucleic acid
 <213> Zea mays

 <400> 864

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 gaactgatac agcaggaact tgaccatgta gttgaagagc ttgagaaaat agcaattgtt 120
 cgtctacttc agcagagggc gataatttca cttatcgga atgtggagca atcgtcnctc 180
 atactagaaa agacgggacg tgtgctgagg aaaagtgggg ttat 224

<210> 865
 <211> 240
 <212> nucleic acid
 <213> Zea mays

 <400> 865

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 cccacccgtt cgcgcactat ctggttccga caacatcata gccttcacca cctcgaggta 120
 caaggatcag cactgatcg tgcgagggtc gggggctggt gcggaagtga ctgcaggagg 180
 cgtcttctgc gacatactgc gcctgtcttc ctaccttggt gcccctcgt agtctgtcaa 240

<210> 866
 <211> 243
 <212> nucleic acid

<213> Zea mays

<400> 866

cttgggatta agagaaattc aggtttggaa agatgttgat ggtgttttaa catgtgaccc 60
 aaatatccat cctaaggcaa aacctgtgcc atacttgaca tttgatgagg cagctgaact 120
 tgcctatfff ggagcgcagg ttttgcattc acaatcaatg cggccagcta gagaaggcga 180
 tgtacctgtg agggtaaga atcttataac cgtcngccc ctggtaccct tatcataaag 240
 cng 243

<210> 867

<211> 286

<212> nucleic acid

<213> Zea mays

<400> 867

gaaggtcata attcttgcaa gggaatctgg tttggggctt gaactttctg atattcctgt 60
 acgaagtctt gtgccggang cttnaagtca tgtacctcag ccgacgaata catgcagaag 120
 ttaccatctt ttgacgagga ttgggcaaga gagaggaaaa atgctgaagc tgcgggagaa 180
 gttctgcggt atgttggagt ggtgnatgtg gtcncaaaaa gggccnaggt gaantnngcc 240
 ctnaaagggg ggccccccnt ttggnnaaaa ntggtcncaa aaaaaa 286

<210> 868

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 868

aagagaagct acaaactgaa gcagaacctg ccaatcttga taagtttggt catcatttgt 60
 ctgagantca cttttttcct aacagagttt tggtagattg cacagcagat acaagtgttg 120
 catctcacta ttatgattgg cttaaagaaag gaatccatgt catcactcca aacaagaaag 180
 caaattcagg gccacttgac aggtatctta aacttagaac tcttcagcgt gcatcatata 240
 ctcatctatt ctatgaagca actgttggtg ctggc 275

<210> 869

<211> 241

<212> nucleic acid

<213> Zea mays

<400> 869

cttttagtgat gttgttgctg aagcaaaaaa agcaggatac acagaaccag atccaagaga 60
 tgatctgtct ggtactgatg ttgccaggaa ggtcataatt cttgcaaggg aatctggttt 120
 ggggctgagc tttctgatat tcctgtacga agtcttgtgc cagangcact taagtcatgt 180
 acctcagccg atgaatacat gcagaagtta ccattctttt acgaggattg ggcaagagag 240
 a 241

<210> 870

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 870

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 tgggtgaata actanttggg caggggtggg agcgacttga cagctacaac cattggcaaa 120
 gccttgggat taagagaaat tcagggttgg aaagatgttg atggtgtttt aacatgtgac 180
 ccaaatatcc atcctaaggc aaaacctgtg ccatacttga cattgatgag gcagctgaac 240
 ttgcctatth ggagcgcagt tttgcatcca caatcaatgc ggccactaga ga 292

<210> 871

<211> 226

<212> nucleic acid

<213> Zea mays

<400> 871

gggaatgcc aggcgggggtt caccttcgtc atgaagttcg gcggctcgtc gctcgcgtca 60
 gccgagcggg tgcgggangt ggccggcctc atccttagct tcctcgacga gatgcctgtc 120
 gccgtcctct ccgcatggg gaagaccacc aataacctn tcctaggacc ttccatttct 180
 agctctcagc gcctgagtca ggggtgctga ttcttcgtta tagtaa 226

<210> 872

<211> 285

<212> nucleic acid

<213> Zea mays

<400> 872

gtcattgatt cttgaaaagg cattcaacgt tctgcgaaga aatgggtgta atgtcnagat 60
gatctcaciaa ggggcgtcca aggtgaacat atccttggtg gtcaatgaca gtgaggcaaa 120
acagtgcgta caagccctcc attcagcgtt ctttgagaat ggattcttgt cggaggtaga 180
gggagctgac gttccccaga acggagcctc tctgaactca aatggcgctg tctatggaaa 240
ctagtcgact ctgcaactgc tttcttcaca atttgtgtac cattg 285

<210> 873

<211> 286

<212> nucleic acid

<213> Zea mays

<400> 873

ccaaggcgtc cgaaatccct gagctcgctg tcatcaagga cctgcatctt aggacggttg 60
atgagctcgg tttagatagg tcgattgttt cagggtttatt ggatgaattg gagcaacttc 120
ttaaaggagt tgctatgatg aaagagctga cccttaggac aagagattac cttgtttcct 180
ttggtgaatg tatgtcgaca agaatatattg ctgcatattt gaacaaaactt gggaaaaaag 240
cacgccatat gatgcattga tatggcttat aaccaccgat gatttc 286

<210> 874

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 874

ttcacaaatg cggatattct tgaagccact taccctgctg ttgcgaagag gctacatgaa 60
gactggatgg atgaccctgc tattcctata gtcactggtt ttcttgggaa gggatgtaaa 120
tcatgtgctg tcaccacttt aaggcagggg tggtagtgc ttgacagcta caaccattgg 180
caaagccttg gggattaaga gaaatccagg ttggaaggat gtagatggtg tgtgacatgt 240
gatcctnata tatatggcaa atgctatanc cctgcctac ttgactttgg atg 293

<210> 875

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 875

actgccaaagt cccttggcta tacagaacca gatccacgtg atgatctagg tggaatggat 60

gtcgctagaa aggcattgat cctggctagg cttcttgggc aacagataag catggagagt 120

atcaatgttg aggtttatat cctagtgaat taggacctga tgcagtgtcc acaaaggact 180

tcctgaaagt ggtttagtgc aattgacaag agcatggaag aaagaatcag agcagcatct 240

cgagaggaaa cgttctcgct acgtct 266

<210> 876

<211> 343

<212> nucleic acid

<213> Zea mays

<400> 876

ggctggttct aggacaaatg cttttaagtg atataggagt agatttgacc cagtggaaaag 60

agaagctaca aactgaagca gaacctgcca atcttgataa gtttgttcat catttgtctg 120

agaatcactt tttccctaac agagttttgg tagattgcac agcagataca agtgttgcat 180

ctcactatta tgattggcta aagaaaggat ccatgtcatc actccaaaca agaagcaant 240

cagggccatt gacaggttct taancttgac tcttcagggg cataattatc atacttctan 300

gagnantgtg gnnngcccca nanatnnccn ngggtnnnan nag 343

<210> 877

<211> 442

<212> nucleic acid

<213> Zea mays

<400> 877

gngatctngn gnantancta ncttgngnan ntgngnggtn cacttnacnn cnannnncat 60

tggnanncn ttgagattaa ganatactcn tgtaggnaa gatgctgatg gtgntggaac 120

atgtgaccga gatatacatc cttaaagctga acctgancca tacttgacat ctngatgaat 180

cagctgaacn ngcctatfff ggagcgcagg aattgcatgc actatnnatg cggtcagngt 240

tgagaaggcg atntacctgt gagggctaata gaattnttat aactcgncta gnctanggta 300

cccttatcac taaaacntgn acgatatgag ttaagantgt tcagactngc attgtncctg 360

aaaatgaata ttnaatgctt gatatatcca gtacagggta tgctttgcca agtaccggat 420

naataanncc cntttaaggg aa

442

<210> 878
<211> 285
<212> nucleic acid
<213> Zea mays

<400> 878

gnncaagacn tgtgccatac ttgacatttg atgaggcagc tgaacttgcc tatnttgag 60
cgcagggtttt gcatccacaa tcaatgcggc cagctagaga aggcgatgta cctgtgaggg 120
ttaagaattc ttataaccgt cganccctg gtacccttat cactaaagca agagatatga 180
gtaagactgt ttgactagc attgttttga aatcaaatac tacaatgctt gatatatgca 240
gcacacgcat gcttgccag tacggttttc tagctaaggt tttt 285

<210> 879
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 879

gncaaaacct gtgccatact tgacatttga tgaggcagct gaacttgcc tatnttgag 60
cgcagggtttt gcatccacaa tcaatgcggc cagctagaga aggcgatgta cctgtgaggg 120
ttaagaattc ttataaccgt cgagccctg gtacccttat cactaaagca agagatatga 180
gtaagactgt ttgactagc attgttttga aatcaaatac tacaagcttg atatagtcag 240
cacacgcatg cttggccagt acggttttct agctaaggtt tt 282

<210> 880
<211> 317
<212> nucleic acid
<213> Zea mays

<400> 880

gaaccagctg atattgatag tttgttcat catttatccg ataatacgt atttccaaac 60
aaagttttgg tggactgcac agcagataca agtgttgcat cccactatta taattgggta 120
aagaagggta ttcatgtcat cacaccaaatac aagaaagcaa attctggggc acttgatcag 180
tacctgaaat tgcggactct gcaacgtgcc tcgtacactc attactttta tgaagcaact 240

gttggtgctg gcctcccaat cattagtacc ttgcgaggtc ttctggagac gggtgacaag 300
 atactgcgta ttgaggg 317

<210> 881
 <211> 315
 <212> nucleic acid
 <213> Zea mays
 <400> 881

ggatcaggtt gctgctctca aggaaaatat gaacattgat ctgcgtgtca ttggaataac 60
 tggctcaagt acaatgcttt tgagcagaca cgggaataga cttaaccag tggaaacaac 120
 ttctaaaaaa ggaagcagaa ccagctgata ttgatagttt tgttcacatc ttatccgata 180
 atcacgtatt tccaaacaaa gttttggtgg actgcacagc agatacaagt gttgcatccc 240
 actattataa ttggttaaag aagggtattc atgtcatcac accaaataag aaagcaaatt 300
 ctgggccact tgate 315

<210> 882
 <211> 288
 <212> nucleic acid
 <213> Zea mays
 <400> 882

aaccagctga tattgatagt tttgttcac atttatccga taatcacgta tttccaaaca 60
 aagttttggt ggactgcaca gcagatacaa gtgttgcac ccactattat aattgggttaa 120
 agaagggtat tcatgtcac acaccaaata agaaagcaaa ttctgggcca cttgatcagt 180
 acctgaaatt gcgactctg caacgtgcct cgtacactca ttacttttat gaagcaactg 240
 ttggtgctgg cctcccaatc atagtacctt gcgaggtctt ctggagac 288

<210> 883
 <211> 292
 <212> nucleic acid
 <213> Zea mays
 <400> 883

ccaaaacgac acttgacgtt ggcacattg gacctggttt aattggtaga acactcctta 60
 accagctcaa ggatcaggtt gctgctctca aggaaaatat gaacattgat ctgcgtgtca 120

cattatcctt gcaagggaaat caggtttgag gcttgaactt tctgatattc ctgtgaagag 180
ccttgtgcc aaaaacttg cagcatgctc gtcagcagat gaattcatgc agaagctacc 240
atcctttgac gaggactggg ccaggcaacg catgatgct 279

<210> 887
<211> 256
<212> nucleic acid
<213> Zea mays

<400> 887

tttantnntt tcgaaggaac aagggcattc agtgntgtcg ttgcngaagc aaaaganntg 60
ggatacactg aacctgacct aaganntgat ctntctggaa ntgntgttgc ccgaaaggtc 120
attntccttg caaggggaatc aggnntgagg cttgaacttt ctggntattc ctgtgaagag 180
ccttgnngctn gaaaacttg cagcagctcg tcagcagaga gttcagcaga agctaccatc 240
ctttgacgag gactgg 256

<210> 888
<211> 322
<212> nucleic acid
<213> Zea mays

<400> 888

gggagactgg atggatgacc ctgctattcc tatagtcact ggttttcntg gncagggatg 60
taaatacatgt gctgtcacca ctttaggcag ggggtgtagt gacttgacag ctacaacccat 120
tggcaaagcc ttgggattaa gagaaatcca ggtttgaag gatgtagatg gtgtgttgac 180
atgtgatcct aatatatatg caaatgctat acccgtgcc tacttgactt ttgatgaggc 240
tgctgagctt gctactttg gtgcacaggt ttgcatccc caatcaatgc gaccggctag 300
ggatggtgat attccagtga ga 322

<210> 889
<211> 105
<212> nucleic acid
<213> Zea mays

<400> 889

gtgatcctaa tatatatgca aatgctatac ccgtgcccta cttgactttt gatgaggctg 60

ctgagcttgc ctactttggt gcacaggtta nacatcccc agtca 105

<210>	890
<211>	279
<212>	nucleic acid
<213>	Zea mays

<400> 890

gtgagtacaa	cattactatt	gtactgaagc	aagaagactg	cgtgcgggct	ctgagagctg	60
cccaactcaag	gntcttcctc	tccaaaacca	caactggctgn	tggcatcatt	ggacctgggn	120
tgattggctg	cacattactt	aatcagctga	aggatcaggc	tgctgtcttc	aaggaaaata	180
tgaacattga	tttgcggtgc	atgggaatag	ctggttctag	gacaatgctt	ttaagtgata	240
taggagtaga	tttgacctag	tggaaagaga	agctacnaa			279

<210>	891
<211>	282
<212>	nucleic acid
<213>	Zea mays

<400> 891

gctatgcaaa	tattaatgta	agggcgatag	ctcaagggttg	cagtgagtac	aacattacta	60
ttgtactgaa	gcaagaagac	tgcgtgcggg	ctctgagagc	tgccactca	aggttcttcc	120
tctccaaaac	cacactggct	gttggcatca	ttggacctgg	gttgattggg	cgcacactac	180
ttaatcagct	gaaggatcag	gctgctgtcc	tcaaggaaaa	tatgaacatt	gtttgcgtgt	240
catgggaata	gctggttcta	ggacaatgct	tttaagtgat	at		282

<210>	892
<211>	282
<212>	nucleic acid
<213>	Zea mays

<400> 892

gttatccatg tttctgaaat cgaagagtgg aatatggtca aaagcctaca tatcaagacg 60
gtggatgaac ttggacttcc aagatctgta atacaagaca tgctagatga actggagcaa 120
ctattgaaag gtatcgctat gatgaaagag ctgacgccta ggaccagtga ctaccttggt 180
tcatttgag aatgcatgtc caccaggatt tttctgctt atttgaacaa aattcgtgtc 240

aaggcacggc agtatgacgc atttgatatt ggtttcatac aa 282

<210>	893
<211>	270
<212>	nucleic acid
<213>	Zea mays

<400> 893

gttatccatg	tttctgaaat	cgaagagtgg	aatatgggtca	aaagcctaca	tatcaagacg	60
gtggatgaac	ttggacttcc	aagatctgta	atacaagaca	tgctagatga	actggagcaa	120
ctattgaaag	gtatcgctat	gatgaaagag	ctgacgccta	ggaccagtga	ctaccttggt	180
tcatttgag	aatgcatgtc	caccaggatt	tttctgctta	tttgaacaaa	attcgtgtca	240
aggcacggca	gtatgacgca	tttgatattg				270

<210>	894
<211>	493
<212>	nucleic acid
<213>	Zea mays

<400> 894

gngnnnnnng gnaantgttn anttngcccg ccagtagcca tccaagaatt cccgggtcga	60
cccacgcgtc cgatcagctt gggctagata gatcaattgt ttgtggtttg ttggatgaac	120
ttgaacaact tcttaaggga attgctatga tgaaagagct gactcttagg acacgagatt	180
atctagtttc atttggtgaa tgcattgtcta caagaatatt tgctgcactt ttaaacaaaa	240
ttggagtaaa agctcggcag tatgatgcat ttgatcttgg ctttataact accgatgatt	300
tcacaaatgc ggatattctg gaagcaactt atcctgctgt tgcaaggagg ctacatgtgg	360
aatggattaa tgatcctgct atacctatag tcaactggctt tcttggcaag ggatggagat	420
ctggtgcaat aactaccttg ggcanggggtg gtaatgactt gacagctaca accattggca	480
aagccttggg aat	493

<210>	895
<211>	281
<212>	nucleic acid
<213>	Zea mays

<400> 895

aaaganctgn ctcttaggac acgagattat ctagttgcag tnggtgaatg catggctaca 60
 agaanattgg cctgcacttt naaacaaaat tggagtaaaa gtctcggcag tatgatgcat 120
 gtnatcttgg cttgataact accgatgatt tcacaaatgc ggntattctg gaagcaactt 180
 atcctgctgt tgcaaggagg cgacatgtgg aatggattaa tgatcctgcn agacctatag 240
 tcantggcnt tcttggcaag ggatggagat ctggtgcaat a 281

<210> 896
 <211> 299
 <212> nucleic acid
 <213> Zea mays

<400> 896

gcgcggggcg ggaggtgact gggggtggcg tcttctgcga catcctgcgg ctggcgtctt 60
 accttggcgc tcttcttag agcttccact tacagttgtc aatttgaaga ggcagatgga 120
 cagtgccttg catgatgcct cgtgttttag gcttattatg gacttgctca aagttgttct 180
 gaataagtgc caggaaataa ggctctatc ttatttaggc ctctatcttc ggggtagctg 240
 actaagtgtt gttatgctac ggaatccaga atctgtgaat tgaaatgaat gtaccgatg 299

<210> 897
 <211> 454
 <212> nucleic acid
 <213> Zea mays

<400> 897

actttacttg gccngccagg tcaagatncc cgggtcgacc cagcntcca ccgcttcngc 60
 cctcncgctt tcgttgtcat gaggagctc accgtggcaa gccgccaccc gggcgccgct 120
 ttctctaccc gccgcgccc cttcttcac ccgcgcgag ccggaaggga ctcgaccttc 180
 cagcggtgct ggcgatggga aaaaactcag gacagctcct tcgggagctc actaaggacc 240
 agccgccttc caagaactgt gcacggagat attttgaaga atttgctcgc accaactgct 300
 ggtgctgttt cagttgagca agctgaagct attgctgacc tcccaaaagg tgaaatgtgg 360
 tcggtgcaca agtttggtgg cacatgcatg gggacctctg agagaattca caatgttgct 420
 gataaaancc ctninggattc cttcgaanaa ggaa 454

<210> 898

<211> 310
 <212> nucleic acid
 <213> Zea mays

 <400> 898

 ccgccctctc gctttcggtg tcatgaggag cctcaccgtg gcaagccgcc acccnggcgc 60
 cgctttctct acccgccgcc gcccccttct tcaccccgcc gcagccggaa aggactcgac 120
 cttccagcgg tgctggcgat gggaaaaaac tcaggacagc tccttcggga antcactaag 180
 gaccanccgc cttccaagaa ntgtgnacgg agatattttg aagnatttgc tcgcaccaac 240
 tgctggtgct gtttcagttg agcaagctga agctattgct gacctccaa aaggtgacat 300
 gtggtcggtg 310

<210> 899
 <211> 276
 <212> nucleic acid
 <213> Zea mays

 <400> 899

 tatgatgtga caggnacttg atcatgtcgt tgaagagctt aaaaaatttg cagttgttca 60
 tctgctgcaa cgtagatcga tcatctcact gatagggat gtacagcgat cgtcattgat 120
 tcttgaaaag gcattcaacg ttctgcgaag aaatgggtgtt aatgtccaga tgatctcaca 180
 aggggcgtcc aaggtgaaca tctcttggt agtcaatgac agtgaggcaa aacagtgcgt 240
 acaagccctc cattcagcgt tctttgagaa tggatc 276

<210> 900
 <211> 240
 <212> nucleic acid
 <213> Zea mays

 <400> 900

 tgaattgac cagcaggaac ttgatcatgt cgttgaagag cttaaaaaat ttgcagttgt 60
 tcattctgtg caacgtagat cgatcatctc actgataggg aatgtacagc gatcgtcatt 120
 gattcttgaa aaggcattca acgttctgcg aagaaatggt gttaatgtcc agatgatctc 180
 acaaggggcg tccaaggtga acatattcctt ggtagtcaat gacagtgagg caaacagtg 240

<210> 901

<211> 279
<212> nucleic acid
<213> Zea mays

<400> 901

ggagcgtcac ccggaagtat gggcccgagg ccgtcgccag cggcgccgta gttgtcgaca 60
acagctccgc gttccggatg gngcccgang tgccgctcgt catccccgag gtcaacccccg 120
aggccatggc gaacgtccgc ctggggcagg gggcgattgt ggcaaaccg aattgctcga 180
ccatcatctg cntcatgggt gccacgccc tccatcgcca cgctaagggtg ttaaggatgg 240
ttgtcagcac ataccaagca gcaatgggtgc ggggtgctgc 279

<210> 902
<211> 254
<212> nucleic acid
<213> Zea mays

<400> 902

cagtgggtag gattcgctcag gacttgtccc tggatgataa ccgaggggtg gacataatttg 60
tgtgtggtga tcagatacgt aaagggtgccg cactcaatgc cgttcagatt gctgaaatgc 120
tgctgaagtg aatgcgactt aaccctcttg tccctccctc cctccctgtc cctaattgct 180
ctgatcaaatt gctggactgt actctgatta gtttgtcctc nattttgggtc gcctgttctg 240
tattctgcgg tgct 254

<210> 903
<211> 496
<212> nucleic acid
<213> Zea mays

<400> 903

ggtnnnnnaa ctccccgc nccggtacgg tncaaaattc ccggnncgac ccacgcgtca 60
caagatgttg tgtacacgga ttggatagat cgtccacaac ctcgattgga ccgtcataag 120
gaaatggggtt tacagtgacg ctgggaagag tgaggggaatg ttccgtgatg gacataagtt 180
ggatttgttt ggacataaca cagtggtagg agctgcgggt ggttcgatta ttantgaaag 240
ttgagtgtcg ccaaagggtt tattgcacaa aaaggaaaac aaagtgtata tgcggagtat 300
ctcgtcacta gtgtgtgtca ctagacagaa tagtgtgttt gtgttgtttg tgtcgtgttt 360

tccaatgaan gacattgacc actagaacta tgtgaaaaaa aaaannagtc aacacaaaga 420
 ntannaaaaa aggnocggccg ctctagagga tncaagctta cgtacccgtc atnacgtcan 480
 aacccttcna aaatgt 496

<210> 904
 <211> 435
 <212> nucleic acid
 <213> Zea mays

<400> 904

gtaactgcga cttgcatacg ggttcctgtg atgcgcgcac atgctgaaag tgtnaagnta 60
 cagtttgaaa agccacttga tgagcgcgta tgctcctcaa gtaggcctta atgagaaactc 120
 aaagagagag ttctgggaag gcoctggagga catgggttagt agtgtgccag ttggcgagaa 180
 gctottcata ggaggagatc tcaatggcca tgatactgca agagaaattt tgagagcagc 240
 tcctggtgtt accattattg atgaccgagc ttccaatcgc ttctctacac ctctggaggt 300
 atcagacaaa gatgacgtan cantgggtag gattcgtcag gacttgtcct ggatggtaac 360
 cganggtttg gacatatttg tgtgttggtg atcagatacg taaangcgcg ggactnaatg 420
 ccgtcaaaat tggtt 435

<210> 905
 <211> 287
 <212> nucleic acid
 <213> Zea mays

<400> 905

atcaccagag tgaattagca gaagaattgg ctgaaaggaa gcatctcttc tgtgcacgcc 60
 cacaaacatt gggagagacc atccgggcaa tggacctaga gacactgggt cctatgagcc 120
 aggggatgcc aaaccagtcg tcaccctgat caacagattt cttggcttcc agttgattga 180
 cgaatctatt cggttggtct caatccatgt ggattagatg gtattgcgtc ggtttaattc 240
 catagcaagt caaaatacat cctaaacca ctcaatccac tccaatc 287

<210> 906
 <211> 307
 <212> nucleic acid
 <213> Zea mays

<400> 906

ccgcgaccga tcgagcgaan ancccttccc gcgcccgcgc cgaaacccta gtcctcttta 60
cgccatggcc acogtgtcgc tcaactccga ggcggtcttc tccaccgagt cggcgggcgc 120
cctgggcttc tgccaccatc ctccgcttcc cgccaaactt cgtacgccag ctacgacca 180
aggcacgacg caactgcagc aacatcgggc tcgcgagat cgtcgccgc gcgtggtccg 240
actgcctcgc tgctcgccgc ctccgggcg gcggatgtca gcgcaattcc taacgctaag 300
gttgcg 307

<210> 907

<211> 265

<212> nucleic acid

<213> Zea mays

<400> 907

aaagtacatt gctggacaca acgatgttat tggaggatgc gtcagtggca gagtgantgg 60
tttccaaagt cntatattat caccatgttg ttggtggtgt ctaaccccga atgctncgta 120
ctancctnnc ncnгааacct ncatctccgt gtgcagtctg aataaacntg cttncggatg 180
gccanttttt agaggaacat ctaaagatgc cngctacta tcctggctgc tagccaaccc 240
tgaacatcca ttgcccagag ncaat 265

<210> 908

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 908

attttcagta aatttggtat cgaacacaca ctgaacatcc ctgcaacttt tggtgttaag 60
angttgtttg tttaggacta taatctgtct agattatata atccaacaaa taatgtgtta 120
tttgctggat tatataatct ggacagatta tattctcaaa caaacacccg ctaattcttc 180
caggggacaa tcggagcgtg gaaacgatgc caaccaggac atactttaca accagaacgt 240
ccaaagtgtg gtgccgtacc attcgag 267

<210> 909

<211> 96

<212> nucleic acid

<213> Zea mays

<400> 909

tacaaatcca tttctcagat gcattgatat tgaacatgta tccaaatatg tgccataggc 60
aaggganngt ngccttngaa tngcngnan ttencg 96

<210> 910

<211> 352

<212> nucleic acid

<213> Zea mays

<400> 910

accaaatcat gcattgtact gaaatgagat gcaattccaa ttctattgtt tggatgtcac 60
tgaattgtac gttctaattc caogcaatac cgaggnnctg angctttgta ttgggagagg 120
atgttctagt tatagtctaa ttctagggaa ttaggcctct gatnccaaat ctcaattcca 180
cgagcaanca aacaacagaa tttagaaaaa ctgattctaa ttcccaattc tgtgtctcaa 240
tatctatatc caaccggggt gttagtatta tgtagtttca ctggctcttc tagttgtgct 300
ttatganatc tttngnacct gaagataaat ttgagttttc aggaaggnac tt 352

<210> 911

<211> 503

<212> nucleic acid

<213> Zea mays

<400> 911

gnnnnnnnnn ggnggaangt ccagccggnc ntaccggtcc canaaattcc cggntcnacc 60
acgcgtccgg agatcttcat caggggaaat tcgcatggta tgggagacta cggcccgtta 120
cctttgaacc ctcgtaggaa gtcttgagaa cccttgcatg cattcgctgt tgtatgctgc 180
catgctgttg aatttagctg tactacacca gatagtata cccgaactca cggagttaat 240
ggaaccggat agaaagctgc ccatgtagtt tgggtgtttgt tgcttacggt gtggatttac 300
tgtagtgact tgtctaattc tccgtttctc cgtttggtta tcgaccatat tggaggcatg 360
gaattaaaat aagttcaata ccaaatcaaa tatagtattg aatgagatg taattcaatt 420
ctattgtttg gatgtcgctg atttgaatt aaaaaanaa nannnaannn nnnnnnnnnn 480
nnnnnnnnnn nnnnnnnnnn nna 503

<210> 912
 <211> 497
 <212> nucleic acid
 <213> Zea mays

<400> 912

ggtttttaan gggggggagg aagggaaagc tttcccggtc cggattcccg ggtcgaccca 60
 cgcgtcccga gttaaaacat ctttttgng gtgggnaaat atggaaccta tttcangtta 120
 nttggcctng nttggatttg ancccgtnng gtggagtaat ctaatactcc gcttgggatg 180
 tcgatattgg agaggatgga attgaattgg ggttgaatac caaatcaggc atgatattga 240
 aatgagatat aatttcaatt ctattgtttg gatgtcactg aattggagtt tggaattatg 300
 cgggtctaatt ctaggcaata ccgagggggg gaggctttgt attggaagag ggggtttcta 360
 gttatagtct aatttcaggg aattgggcct ctgattccaa atctcaattc cacatgcaac 420
 gaatcaacat aatttangaa agctgattcc aattctcaat tctgtgattc aatctctaca 480
 tccaaacggg gtataag 497

<210> 913
 <211> 548
 <212> nucleic acid
 <213> Zea mays

<400> 913

gnnnngcgca gagtttgttn gggngnnggg gggaaggang natntgctnn tccggtccgg 60
 aattcccggg tcgaccacg cgtccgcgcg agcggcccc tttttttttt tttttttttt 120
 ttttttttta ctaaaaccta tngnctgang ntcaggctaa tgacattttg tctttacaat 180
 actgtgatgc aataaaactg ggcaaaccat tgataccacc gtatacagta actcaatacc 240
 atccaattct atgaggatta aaaatgtatt ggaatggatt ggtatgtatt ttgacttgct 300
 ataaatttaa accaactnaa taccatccaa ttcacttaga ttaacggtga aacgaacaag 360
 gcctaagggg agtggttaaca gtctaacatc attgtacaac agcctttccg aagcccatag 420
 acattgtgag acaagatccg taaaccatga atcgttcttc acttaggatg accatnttag 480
 ctggtcacga aatgggtaccg ggaatcacia atntttnaac gtcccaaang aacgatcctg 540
 gangctgt 548

<210> 914
 <211> 463
 <212> nucleic acid
 <213> Zea mays

<400> 914

cataaataag tcgtgttttg aagggttctn ntgaataaga taaaggctta gatacattna 60
 gntgtatgaa cgggttnttt ttaggtgtat ctagtgtttg gagaaccgca acatgatgtc 120
 tatgttgtac atgacctaag tgcatttctc taccaccagc actcttgccc aagcatggta 180
 gaccatttgt gtttaatgtt cattacatgc taataaaaga tactggtggg aagggttaagg 240
 ttatgtttta ggcacaccaa acaagaagga gtctaangaa attaagnnac caaggctctc 300
 agtngagtaa gggggtagtc tttccattaa aattgactgg ctaaaattta agctcgggct 360
 cttatgngng ttngngnaac ctatgagaac cgacngnaag gnanttgggn aatcagtnag 420
 tanttgaac naaacggtnt ggtaaagtga acccttttcc cgt 463

<210> 915
 <211> 267
 <212> nucleic acid
 <213> Zea mays

<400> 915

agctgaagga tttcttcaag cagtaggtgt gaagacctga agatcagctt tatctctctt 60
 ttgttctctg tagggccttc tcggttattc caattccatg tggattgaag tgtattaaaa 120
 tggattttga cttgttatgg atttaaatac acttaatacc actcaatcca catggattat 180
 tggatgaaaca aacaagccct agtcgaaatt cttgttaagt tggtagtttg tcagaccoga 240
 aatagagttt tttgggtttc gaatgga 267

<210> 916
 <211> 284
 <212> nucleic acid
 <213> Zea mays

<400> 916

attttgaggc atggcagaca actgaggtcg aaatagcgtg gtagcttgga tctgcacatc 60
 tatatgattg agggcctggg gtttaggatt ataatctgcc cagattatat aacctatctt 120

atTTTgaact aagtgttagt tcaaaataag ttagattata taatatggac aaattatatt 180
 cccaatcaaa caggctatga ttggacttac tgatacccat tgagtagaac tggatcttgg 240
 acagtntnCG gggaaaacgg atggacctta attgaaaatt tgcg 284

<210> 917
 <211> 247
 <212> nucleic acid
 <213> Zea mays
 <400> 917

ctgaggTcga aatagcgtgg tagcttggat ctgcacatct atatgattga gggcctggTg 60
 tttaggatta taacctgcc agattatata acctatctta tttgaactaa gtgttagttc 120
 aaaataagtt agattatata atctggacaa attatatctc caatcaaaca ggctatgatt 180
 ggacttactg ataccattt gagtagaact gatcttgaca ggtatccgag aaaacgatga 240
 ccttaat 247

<210> 918
 <211> 254
 <212> nucleic acid
 <213> Zea mays
 <400> 918

caactggagg tcgaaatagc gtggttagctt ggatctgcac atctatatga ttganggcct 60
 ggtgttttagg attataatct gccagatta tataacctat cttattttga actaagtgtt 120
 agttcanaat aagttagatt atataatatg gacaaattat attcccaatc aaacaggcta 180
 tgattggact tactgatacc catttgagta gaactgatct tgacaggtat ccgagaaaac 240
 gntgacctta attg 254

<210> 919
 <211> 508
 <212> nucleic acid
 <213> Zea mays
 <400> 919

agangttttt nggntggggn ggnnganaag ttaccggTc cggattcccg ggtcgaccca 60
 cgcgtccgct gggaagaagg atagccacgg atgctatcac cagcccggtg gtgaacacgt 120

cggcctactg gttcaacaac tcgcaagagc taatcgactt taaggagggg aggcattgcta 180
gcttcgagta tgggaggtat gggaacccga ccacggaggc attagagaag aagatgagcg 240
cactggagaa agcagagtcc accgtgtttg tggcgtcagg gatgtatgca gctgtggcta 300
tgctcagcgc acttgteccct gctgggtgggc acattgtgac caccacggat tgctaccgca 360
agacaaggat ttacatggaa aatgagctcc ctaagagggg aatttcgatg actgtcatta 420
ggcctgctga catggatgct ctccaaaatg cgttggacaa caataatgta tctcttttct 480
tcacggagac tntacaaaat ccatttct 508

<210> 920
<211> 316
<212> nucleic acid
<213> Zea mays

<400> 920
ggagactccc acaaattccat ttctcagatg cattgatatt gaacatgtat caaatatgtg 60
ccatagcaag ggagcggttg tttgtatcga cagtactttt gcctccccta tcaatcagaa 120
ggcactgact ttaggcgctg acctagtatt tcattctgca acaaagtaca ttgctggaca 180
caacgatgtt attggaggat gcgtcagtg cagagatgag ttggtttcca aagtccgtat 240
ttatcaccat gtggttggtg gtgttctaaa cccgaataac actgctcttc ggatggccca 300
gttttttagag gaacat 316

<210> 921
<211> 300
<212> nucleic acid
<213> Zea mays

<400> 921
ttcgatgact gtcattaggc ctgctgacat ggatgctcta caaatgcgt tggacaacaa 60
taatgtatct cttttcttca cggagactcc cacaaatcca tttctcagat gcattgatat 120
tgaacatgta tcaaatatgt gccatagcaa gggagcggtg ctttgtatcg acagtacttt 180
tgctcccct atcaatcaga aggcactgac ttaggcgct gacctagtta ttcattctgc 240
aacaagtac attgctggac acaacgatgt tattggagga tgcgtcagtg gcagagatga 300

<210> 922

<211> 271
 <212> nucleic acid
 <213> Zea mays

<400> 922

cttggtccgg ctggtgggca cattgtgacc accacggatt gctaccggaa aacaaggatt 60
 tacatggaaa ctgagctccc caagagggga atttcgatga ctgtcattag gctgtctgac 120
 atggatgctc tacaaaatgc gttggacaac aataatgtat ctcttttctt cacggagact 180
 cccacaaatc cattttctcag atgcattgat attgaacatg tatcaaatat gtgccatagc 240
 aaggggagcgt tgctttgtat cgacagtact t 271

<210> 923
 <211> 266
 <212> nucleic acid
 <213> Zea mays

<400> 923

cacaaatcca tttctcagat gcattgatat tgaacatgta tcaaatatgt gccatagcaa 60
 gggagcgttg ctttgtatcg acagtacttt tgccctccct atcaatcaga aggcactgac 120
 tttaggcgtc gacctagtta ttcattctgc aacaaagtac attgctggac acaacgatgt 180
 tattggagga tgcgtcagtg gcagagatga gttggtttcc aaagtccgta tttatcacca 240
 tgtggttggt ggtgttctaa acccga 266

<210> 924
 <211> 263
 <212> nucleic acid
 <213> Zea mays

<400> 924

tgctcagtg acattgttccg gctggtgggc acattgtgac cacacggatt gctaccggaa 60
 aacaaggatt tacatggaaa ctgagctccc caagagggga atttcgatga ctgtcattag 120
 gctgtctgac atggatgctc tacaaaatgc gttggacaac aataatgtat ctcttttctt 180
 cacggagact cccacaaatc cattttctcag atgcattgat attgaacatg tatcaaatat 240
 gtgccatagc aaggggagcgt tgc 263

<210> 925

<211> 238
<212> nucleic acid
<213> Zea mays

<400> 925

gcgtaacctg ctcggtccg acgccagcct cgccgtccac gcgggggaga ggctgggaag 60
aaggatagcc acggatgcta tcaccacgcc ggtagtgaac acgtcggnct actggttcaa 120
caactcgcaa gagctaatacg actttaagga ggggaggcat gctagcttcg agtatgggag 180
gtatgggaac ccgaccacgg aggcattaga gaagaagatg agcgactgg agaaagca 238

<210> 926
<211> 332
<212> nucleic acid
<213> Zea mays

<400> 926

gccgcaactg cagcaacatc ggcgtcgcgc agatcgtcgc cgccgctgg tccgactgcc 60
ccgccgcctc cgcggcgcgg aggtcagcgc aattcccaac gctaagggtg cgcaaaccga 120
ccgccgtcgt cttggccgag cgtaacctgc tcggctccga cgccagcctc gccgtccacg 180
cggggganag gctgggaaga aggatcgcca cggtatgcgat caccacaccg gtagtgaaca 240
cgtcggccta ctggttcaac aactcgcaag agctaatacg cttaaggag gggaggcatg 300
ctagcttcga gtatgggagg tatgggaacc cg 332

<210> 927
<211> 226
<212> nucleic acid
<213> Zea mays

<400> 927

ctatgctcag tgcacttggt ccggctggtg ggcacattgt gaccaccacg gattgctacc 60
ggaaaacaag gatttacatg gaaactgagc tcccaagag gggaatttcg atgactgtca 120
ttaggcctgc tgacatggat gctctacaaa atgcgttga caacaataat gtancccttt 180
tcttcacgga gactcccaca aatccatttc tcagatgcat tgatat 226

<210> 928
<211> 228
<212> nucleic acid

<213> Zea mays

<400> 928

gatgactgtc attaggcctg ctgacatgga tgctctacaa aatgcgttgg acaacaataa 60
tgtatctctt ttcttcaagg agactccac aaatccattt ctcagatgca ttgatattga 120
acatgtatca aatatgtgcc atagcaaggg agcgttgctt tgtatcgaca gtacttttgc 180
ctcccctatc aatcagangg catgacttta ggcgtgacc tagttatt 228

<210> 929

<211> 501

<212> nucleic acid

<213> Zea mays

<400> 929

gnnagnnaga ggttgatacg gaacgggggn ttaatggaat gctcgtagcg gtccggaatt 60
cccgggtcga cccacgcgtc cgggttattg gaggatgcgt cagtggcaga gatgagttag 120
tttccaaagt tcgnatttac caccatgtag ttggtggtgt tctaaacccg aatgctgcgt 180
acottatcct tcgaggtatg aagacactgc atctccgtgt gcaatgtcag aacgacactg 240
ctcttcggat ggcccagttt ttagaggagc atccaaagat tgctcgtgtc tactatcctg 300
gcttgccaag tcaccctgaa catcacattg ccaagagtca aatgactggc tttggcggtg 360
ttgttagttt tgaggttgct ggagactttg atgctacgag gaaattcatt gattctgtta 420
aaatacccta tcatgcgcct tcttttgaag ctgtgagagc ataattgatc agcctgccat 480
catgtcctac tgggattcaa a 501

<210> 930

<211> 489

<212> nucleic acid

<213> Zea mays

<400> 930

aacgttacgc cggncctag cctccaaaa ttcangggtc gacccacgcg tccggacaac 60
ctgatcaggt tcagcattgg agtggaggat tttgaggacc ttaagaatga tcttgttcag 120
gccctcgaga agatataagc tccaatccat ttgcattaat aaagttatga ggtgatgggt 180
gtctggatct tgtagagatt tgtgatgata catgagctgt tgactgcgat taagttctct 240

tgtgcttatt ttattctttg aattcaataa gctgtggttg ctgcatcttg catgtagtgt 300
 tggacttaat ttacttgctc aaattacaat agtttggtcc catgtacttg tgtccgcgcc 360
 tccgcggttg tgccccaag tgtttttgtg tcatttgctg tgtaatgtga gattaagggc 420
 acgatgtcct taatcaccat aataaatcca agccttaaat gtaaggattc tgggttgga 480
 atgaanaac 489

<210> 931
 <211> 340
 <212> nucleic acid
 <213> Zea mays

<400> 931

tttaaaatac cctatcatgc gccttctttt ggaggctgtg agagcataat tgatcagcct 60
 gccatcatgt cctactggga ttcaaaggag cagcgggaca tctacgggat caaggacaac 120
 ctgatcaggt tcagcattgg tgtggaggat ttcgaggatc ttaagaacga tctcgtgcag 180
 gccctogaga agatctaagc actctaata gtttgtattg acaaaatcat gaggtgatgg 240
 ctgtcttgga tcttgtcaag atctgtgaca atgatatgag ctgatgactg cgaataagtt 300
 ctcttttgc tatttatccg tcaaatccat aagctgtgat 340

<210> 932
 <211> 303
 <212> nucleic acid
 <213> Zea mays

<400> 932

ctgcatctcc gtgtgcaatg tcagaacgac actgctcttc ggatggcca gtttttagag 60
 gagcatccaa agattgctcg tgtctactat cctggcttgc caagtcacc tgaacatcac 120
 attgccaaga gtcaaagac tggctttggc ggtgttgta gttttgaggt tgctggagac 180
 tttgatgcta cgaggaaatt cattgattct gttaaaatac cctatcatgc gccttctttt 240
 ggaggctgtg agagcataat tgatcagcct gccatcatgt cctactggga ttcaaaggag 300
 cag 303

<210> 933
 <211> 299
 <212> nucleic acid

<213> Zea mays

<400> 933

tttacattta aggcattgat ttattatggt gattaaggac atcgtgcctt aatctcacac 60
 tacacagcaa atgacacaaa aacactaggg gcacaaccgc ggaggcgcgg acacaagtac 120
 atgggaccaa actattgtaa ttgacaagt aaaattaagt ccaacaactac atgcaagatg 180
 cagcaaccac agcttattga attcaaagaa taaaataagc acaagagaac ttaatcgag 240
 tcaacagctc atgtatcatc acaaattctc acaagatcca gacaaccatc acctcataa 299

<210> 934

<211> 345

<212> nucleic acid

<213> Zea mays

<400> 934

gggaacatct aaagattgcc cgtgtctact atcctggctt gcctagccac cctgaacatc 60
 acattgccaa gagtcaaattg actggctttg gtggtgtggt tagttttgag gttgctggag 120
 actttgatgc cacaaggaaa ttcattgatt ctgttaagat accctatcat gccccttctt 180
 tnggaggctg tgagagcata attgatcagc ctgctatcat gtcctactgg gattcaaagg 240
 agcagaggga catctatggg atcaaggaca acctgatcag gttcagcatt ggagtggagg 300
 ttttgaggac cttaagaatg atcttgttca ggccctcgag aagat 345

<210> 935

<211> 274

<212> nucleic acid

<213> Zea mays

<400> 935

agtcaaata ctggctttgg cgggtgtgtt agttttgagg ttgctggaga ctttgatgct 60
 acgaggaaat tcattgattc tgttaaaata ccctatcatg cgccttcttt tggaggctgt 120
 gagagcataa ttgatcagcc tgccatcatg tcctactggg attcaaagga gcagcgggac 180
 atctacggga tcaaggacaa cctgatcagg ttcagcattg gtgtggagga tttcgaggat 240
 cttaagaacg atctcgtgca ggccctcgag aaga 274

<210> 936

<211> 306
 <212> nucleic acid
 <213> Zea mays

 <400> 936

 gttggtttcc aaagtccgta tttatcacca tgtggttggt ggtgtttctaa acccgaatgc 60
 tgcgtacctt atccttcgag gcatgaaaac acttcattctc cgtgtgcaat gtcagaataa 120
 nactgctctt cggatggccc agttttttaga ggaacatcta aagattgccc gtgtctacta 180
 tcctggcttg cctagccacc ctgaacatca cattgccaaag agtcaaata ga ctggctttgg 240
 tgggtgtggtt agtttttnagg ttgctggaga ctttgatgcc acaaggaaat tcatcgattc 300
 tgttaa 306

<210> 937
 <211> 265
 <212> nucleic acid
 <213> Zea mays

 <400> 937

 ccaccatgta gttggtggtg ttctaaaccc gaatgctgcg taccttatcc ttcgaggtat 60
 gaagacactg catctccgtg tgcaatgtca gaacgacact gctcttcgga tggcccagtt 120
 tttagaggag catccaaaga ttgctcgtgt ctactatcct ggcttgccaa gtcaccctga 180
 acatcacatt gccaaagatc aaatgactgg ctttggcggg gttgttagtt ttgagggttc 240
 tggagacttt gatgctacga ggaaa 265

<210> 938
 <211> 322
 <212> nucleic acid
 <213> Zea mays

 <400> 938

 ctattggttg tgtggttagt tttgaggttg ctggagactt tgatgccaca aggaaattca 60
 tcgattctgt taagatacnc tatcatgccc cttctttcgg aggctgtgag agcataattg 120
 atcagcctgc tatcatgtcc tactgggatt caaaggagca gagggacatc tatgggatca 180
 aggacaacct gatcaggttc agcattggag tggaggattt tgaggacctt aagaatgatc 240
 ttgttcaggc cctcgagaag atataagctc caatccattt gcattaataa agttatgagg 300

tgatggttgt ctggatcttg ta

322

<210> 939
<211> 283
<212> nucleic acid
<213> Zea mays

<400> 939

cattgccaaag agtcaaata ctggccttgg cgggtgttgtt agttttgagg ttgctggaga 60

ctttgatgct tacgagggaa attcattgat tctgttaaaa taccctatca tgcgccttct 120

tttgaggagct gtgagagcat aattgatcag cctgccatca tgtcctactg ggattcaaag 180

gagcagcggg acatctacgg gatcaaggac aacctgatca ggctcagcat ggtgtggagg 240

attcgaggat cttaagaacg atctcgtgca ggccctcgag aag 283

<210> 940
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 940

ctttagatgt tcctctaaag attgcccgtg tctactatcc tggcttgcct agccaccctg 60

aacatcacat tgccaagagt caaatgactg gctttggtgg tgtggttagt tttgagggtg 120

ctggagactt tgatgccaca aggaaattca tcgattctgt taagataccc tatcatgccc 180

cttctttcgg aggctgtgag agcataattg atcagcctgc tatcatgtcc tactgggatt 240

caaaggagca gagggacatc tatgggatca aggacaacct ga 282

<210> 941
<211> 315
<212> nucleic acid
<213> Zea mays

<400> 941

gggattcaaa ggagcagagg gacatctatg ggatcaagga caacctgatc aggttcagca 60

ttggagtgga ggattttgag gaccttaaga atgatcttgt tcangccctc gagaagatat 120

aagctccaat ccatttgcac taataaagtt atgangtgat ggttgtctgg atcttgtaga 180

gatttgtgat gatacatnag ctntttactg ngaataagtt ctcttgngct taatttantic 240

tttgnattca ataantgtn ggttgcttgg aaaaaaaaaa aannntnnnn nnnnnntnnn 300
 tttttnnnnt nnnnt 315

<210> 942
 <211> 261
 <212> nucleic acid
 <213> Zea mays
 <400> 942

atttatcacc atgtggttgg tgggtgttcta aaccogaatg ctgcgtacct tatccttoga 60
 ggcatgaaaa cacttcatct ccgtgtgcaa tgtcagaata acaactgctct tcggatggcc 120
 cagtttttag aggaacatct aaagattgcc cgtgtctact atcctggctt gcctagccac 180
 cctgaacatc acattgcaa gagtcaaag actggctttg gtgggtgtgg tagttttgag 240
 gttgctggag actttgatgc c 261

<210> 943
 <211> 277
 <212> nucleic acid
 <213> Zea mays
 <400> 943

ttagggcgct gacctagtta ttcattctgc aacaaagtac attgctggnc acaacgatgt 60
 tattggagga tgcgtcagt gcanagatga gttggnttcc naagncgtat ttatcaccat 120
 gtggttgggt gtgtcctaaa cccgaatgct gcgtacctta tccttcgagg catgaaaaca 180
 cttcatctcc gtgtgcaatg tnagaataac actgctcttc ggatggccca gtttttagag 240
 gaacatctaa agattgcccg tgtctactat cctggct 277

<210> 944
 <211> 250
 <212> nucleic acid
 <213> Zea mays
 <400> 944

tgcgtcagt gacagagatga gttagtttcc aaagttcgta tttaccacca ttagtttgg 60
 ggtgttctaa acccgaatgc tgcgtacctt atccttcgag gtatgaagac actgcatctc 120
 cgtgtgcaat gtcagaacga cactgctctt cggatggccc nngttttaga ggagnatcca 180

aagatttcgc gnggtcaagt atccgggctt gccaaagtcac cctgaacatc acattgccaa 240
gagtcaaattg 250

<210> 945
<211> 303
<212> nucleic acid
<213> Zea mays

<400> 945

aggaaatcat cgattctgtt aagaaaccct atcatgcccc ttctttcgga ggctgtgaga 60
gcataattga cagcctgcta tcatgtctac tgggattcaa aggagcagag gggcatctat 120
gggatcaagg acaaccttga caggtcagca ttggagtggg ggattttgag gaccttaaga 180
atgatcttgn tcaggccctc gagaagatat aagctccaat ccatttgcac taataaagggt 240
atgaggtgat ggggtggctgg atcttgagag attgngatga tncatgagct gttgctgca 300
tta 303

<210> 946
<211> 239
<212> nucleic acid
<213> Zea mays

<400> 946

ctatcatgcc ctttctttcg gaggtgtga gaggcataatt gatcagcctg ctatcatgtc 60
ctactgggat tcaaaggagc agaggacat ctatgggatc aaggacaacc tgatcagggtt 120
cagcattgga gtggaggatt ttgaggacct taagaatgat cttgttcagg ccctcgagaa 180
gatataagct ccaatccatt tgcattaata aagttatgag gtgatgggtg ctggatctt 239

<210> 947
<211> 248
<212> nucleic acid
<213> Zea mays

<400> 947

atttatcacc atgtggttgg tgggtgttcta aaccggaatg ctggttacct tacccttga 60
ggcatgaaaa cacttcatct ccgtgtgcaa tgcagaata aactgctct tcggatggcc 120
cagtttttag aggaacatct aaagattgcc cgtgtctact atccnggctt gctagccac 180

ccngaacatc acattgccaa gagtcaaag actggccttg gtgggtgtgg tagttttgag 240
gttgctgg 248

<210> 948
<211> 223
<212> nucleic acid
<213> Zea mays

<400> 948

gatcagcctg ctatcatgtc ctactgggat tcaaaggagc agagggacat ctatgggatc 60
aaggacaacc tgatcagggt cagcattgga gtggaggatt ttgaggacct taagaatgat 120
cttggtcagg cctcgagaa gatataagct ccaatccatt tgcattaata aagttatgag 180
gtgatgggtg tctggatctt gtagagattt gtgatgatac atg 223

<210> 949
<211> 236
<212> nucleic acid
<213> Zea mays

<400> 949

ggagactttg atgccacaag gaaattcatc gattctgtta agatacccta tcatgcccct 60
tctttcggag gctgtgagag cataattgat cagcctgcta tcatgtccta ctgggattca 120
aaggagcaga gggacatcta tgggatcaag gacaacctga tcaggttcag cattggagtg 180
gaggattttg aggaccttaa gaatgatctt gttcaggccc tcgagaagat ataagc 236

<210> 950
<211> 181
<212> nucleic acid
<213> Zea mays

<400> 950

gctctgtggc ttataaacag ggattcaaag gagcagaggg acatctatng gatcaaggac 60
aacctgatca gggtcagcat tggatggagg attttgagga ccttaagaat gatcttggtc 120
aggccctcga gaagatataa gctccaatcc atttgcatta ataaagttat gaggtgatgg 180
t 181

<210> 951

<211> 178
 <212> nucleic acid
 <213> Zea mays

<400> 951

cacaacgatg ttattggagg atgcgtcagt ggcagagatg agttggtttc caaagtcctg 60
 atttatcacc atgtgggttg tgggtgttcta aaccggaatg ctgcgtacct tatecttcga 120
 ggcatgaaaa cacttcatct ccgtgtgcaa tgtcagaata aactgtctct tcggangg 178

<210> 952
 <211> 308
 <212> nucleic acid
 <213> Zea mays

<400> 952

ancgaanacn gctccccgag ccgcgcgcaa aaccctagct tctctactc catggccact 60
 gtctcgctca cccgcgaggc tgtcttctcc acggagtcgg gtggcgccct ggctctgct 120
 accatcctcc gctttccgcc aaactttgtc cgccagctta gcaccaaggc acgccgcaac 180
 tgcagcaaca tggcgctgc gcagatcgtc gccgcgcgt ggtccgactg cncgcngcc 240
 tccgcggcgc ggaggtcagc gcaattccca acgctaaggt tgcgcaaccg tccgcngtcg 300
 tcttggcc 308

<210> 953
 <211> 269
 <212> nucleic acid
 <213> Zea mays

<400> 953

gccacctccc cgaaagctgc cgcgaccgat cgagcgaagn ncggctcccc gcncccgccg 60
 ccaaaacctt agcttctctt actccatggc cactgtctcg ctacccccgc aggtgtctt 120
 ctccacggan tccggtggcg cctggcctc tgctaccatc ctccgctttc cgccaaactt 180
 tgtccgccag cttagcacca aggcacgccg caactgcagc aacatcgccg tcgcgcagat 240
 cgtcgccgcc gctggtccg actgccccg 269

<210> 954
 <211> 276
 <212> nucleic acid

<213> Zea mays

<400> 954

ctgcaaaatg ctgaagggtc gggctctggca ccttttgact gctggctttg cttgagggga 60
 atcaaaacca tggctctgcg ggtggagaaa caacaggcta atgccagaa gattgctgaa 120
 ttcctggcgt ctaccccgag ggtcaagcaa gtaaaactacg ctgggcttcc tgaccatcct 180
 gggcgagctt tacactattc ccaggcnaag ggancgggct ctgttctcag ttttcccacc 240
 ggntaaatgg ccntcnaaa gnagggtggg ggaacc 276

<210> 955

<211> 440

<212> nucleic acid

<213> Zea mays

<400> 955

cgagagtaga cgatcagaac caccatata gtctcacca gcaagttgtg ccaagaaatg 60
 gaatagttgt aaaacgagta gatacaacga aaattagtga tgtggtgtct gcaattggac 120
 cctccactag actggtttgg ctcgaaagtc ccacgaacct tcgtcagcaa attactgaca 180
 ttaagacaat ctgagagata gcgcattctc atgggtgtct tgttttggtt gacaacagca 240
 tcatgtctcc agtgtctctc cgtcctatag aactgggagc tgatatcgtg atgcactcgg 300
 ctaccaaatt tataacggga catagtgatc ttaagggctg ggaaatcttg cagtgaagg 360
 tganaatttg ggctaaaaga aggtaagggt ttctgcaaaa tgctgaaag ggnccgggtcc 420
 gggnaccttt tgactggctg 440

<210> 956

<211> 283

<212> nucleic acid

<213> Zea mays

<400> 956

cgtcgtggag accaccaagt acttcagcgt aacagtcagc ttcgggagcg tgaagtcct 60
 catcagcctg ccgtgcttca tgtcccacgc atcaatcctt gctcgggtcc gcgaggagcg 120
 tggcctaacc gacgacctcg tccggatata ggtcggcatc gaggatgtcg aggacctcat 180
 cgccgatctg gaccgcgcgc tcagaactgg cccggtgtag acatcgccga tccttaggtc 240

atgtcaagct atcttttgat gattcattgg ttgactgctt gcg 283

<210> 957
 <211> 340
 <212> nucleic acid
 <213> Zea mays

<400> 957

caagtacttc agcgtaacag tcagcttcgg gagcgtgaag tccctcatca gcntgcoctg 60
 cttcatgten caagcatcaa tccctgcctc ggtcccgcan gagcgtggcc taaccgacga 120
 cctcgctccg atatcggtcg gcatcgagga tgcgaggac ctcatcgccg atctggaccg 180
 cgcgctcaga actggcccgg tgtagacatc gccgatcctt aggtcatgtc aagctatctt 240
 ttgatgattc attggttgac tgcttgctg atgataataa tgggaatgtt gcttgatac 300
 gccgctgcta naaaaaaact gngtcnagnn acngggnnnc 340

<210> 958
 <211> 257
 <212> nucleic acid
 <213> Zea mays

<400> 958

gcattctcat ggtgctcttg ttttggttga caacagcatc atgtctccag tgctctcccg 60
 tcctatagaa ctgggagctg atatcgtgat gcactcggct accaaattta tagcgggaca 120
 tagtgatctt atggctggaa ttcttgcaat gaagggtgag agtttggtta aagaggtagg 180
 gtttctgcaa aangctgaag ggtcgggtct ggcacctttt gactgctggc tttgcttgag 240
 gggaatcaaa accatgg 257

<210> 959
 <211> 521
 <212> nucleic acid
 <213> Zea mays

<400> 959

ggnnnagggg cttnngtntn tgacntntac ttnntggcat gccgtaccgg tccggaattc 60
 ccgggtcgac ccacgcgtcc ggggtgctgc aattggacct tccactagac tggtttggct 120
 cgaaagtccc acgaaccctc gtcagcaa at tactgacatt aagacaatct cagagatagc 180

acatttctcat ggtgctcttg ttttggttga caacagcatc atgtctccag tgctctcccg 240
tcttatagaa ctgggagctg atatcgtgat gcactcggct accaaattta tagcgggaca 300
tagtgatctt atggctggaa ttcttgcaagt gaaggggtgaa aagtttggtt aaagaggtan 360
ggtttctgca aaatgctgaa ngggtcgggt ctggcacctt ttgactgntg gctttgcttg 420
aagggaatca aaaccatggc tntgcnggtg gagaaacaac aggctaattgc ccagaagatt 480
gctgaatttc tggggtnnta cccganggtc aaagcaagta a 521

<210> 960
<211> 265
<212> nucleic acid
<213> Zea mays

<400> 960
atcgtgatgc actcggctac caaatttata gcgggacata gtgatcttat ggctggaatt 60
cttgcaagtga agggtagagag tttggctaaa gaggtagggt ttctgcaaaa tgctgaaggg 120
tcgggtcttg caccttttga ctgctggctt tgcttgaggg gaatcaaaac catggctctg 180
cggttgagga aacaacagggc taatgccag aagattgctg aattcctggc gtctcaccgg 240
agggtaagc aagtaaacta cgctg 265

<210> 961
<211> 148
<212> nucleic acid
<213> Zea mays

<400> 961
atatcgtgat gcactcggct accaaattta tagcgggaca tagtgatctt atggctggaa 60
ttcttgcaagt gaagggtagag agtttggtta aagaggtagg gtttctgcaa aatgctgaag 120
ggtcgggtct ggcacctttt gactgctg 148

<210> 962
<211> 453
<212> nucleic acid
<213> Zea mays

<400> 962
ggctggggcg tcgctgacgg tgatcgagg cgctccggc ggctccgaac gagatctgag 60

cgctccgca gtctccgtgg aggccctgga ctccgtcgcc tccgattctg acttagagac 120
gaaggagccc agtgtgtcga cgatgctgac gagcttcgag aactcggtcg acaagtatgg 180
ggctctgagc acaccgctgt accagaccgc cacctttaag cagccttcag ctacagatta 240
tggaacttat gattacacta gaagtggtaa cctactcgt gatgttctcc agagcctcat 300
ggctaagctt gagaaagcag atcaagcatt ctgcttcacc agcgggatgg cggcggttagc 360
tgcagtaaca caccttcttc angctggaca agaaatagtt gctggtgang acatatatgg 420
tggttctgat cggctctactt ttgcaagttg tgc 453

<210> 963
<211> 230
<212> nucleic acid
<213> Zea mays

<400> 963
cagcgccctcc gcagtctccg tggaggccct ggactccatc gcctccnatt ctgacttaga 60
gacgaaggag ccagtggtgt cgacgatgcn nacgagnttc gagaactcgt tcgacaagta 120
tggggctctg agcacaccgg tgtaccagac cgccaccttt aagcagcctt cagntacaga 180
ttatggaact tatgattaca ctagaagtgg taaccctact cgtgatgntc 230

<210> 964
<211> 212
<212> nucleic acid
<213> Zea mays

<400> 964
ccgtgctgga caccaacatc ctctgggtga accccgactg cggantcaag acccgaaagt 60
acacggaggt caagcccgcc ctgaccaaca tggctccgc tgctaactca ttgcaccca 120
gctngccagc gccaatgagc ncttttttgc tttntcgttg ggaggnggcn tnttanttct 180
nttttttnaa nanagggccccc cccctttttt nt 212

<210> 965
<211> 275
<212> nucleic acid
<213> Zea mays

<400> 965

ggnccaaaca ccaagttctc ctatgcttct cacaaggctg ttaacgaata taaggaggnt 60
 aaggngctcn gtgttganen cgtgcnagta nttgttngnc tagtnntgta gctgtngatc 120
 tcaaagcctg ccaagggatg ggagaacgga tncccnctgg cttgccctta ctaggacgat 180
 nctccnctgc tacaaggaag tgnntggenn aattnaaatg nagnnnnggg ttccaggatt 240
 canttttang agcccgattg tgnngcnca tgaat 275

<210> 966
 <211> 254
 <212> nucleic acid
 <213> Zea mays

<400> 966

caagacccta acatctctga gcagtgtgac tgcttatggt tttgatcttg tccgtgggtg 60
 ggcttgtcac gagtgtggt ttccctgctg gaaagtacct ctttgcctgn gttgtggatg 120
 gacgcaacat ctgggtgat gatcttgcta catcnctnag naccncaagt cnnntgantn 180
 tgnntttggg aagnaangct tgnngancaa ctgctgcgcg tnaggcacng cgcntgnnnc 240
 tngtaaanaa caat 254

<210> 967
 <211> 313
 <212> nucleic acid
 <213> Zea mays

<400> 967

cgatatggnt gactacttcg gtgagcagct ctccggtttt gcgttactg ccaattgctg 60
 ggtgcagtct atnggtcaag gtgctgcaag cctccgatca tctacggtga cgtgagccgg 120
 cccaacccca tgaccgtttc tngtcgaaga cggcacagag catgacgtct cgcccaatga 180
 aggggatgct gactggncat cacatctcac tgtccttcgt cggattnaca gcgaggttgn 240
 gantgtacag angtcttgga tcaagaagga gngaggncct gagctgtgta tccgtaccaa 300
 cgcgagtnnt gaa 313

<210> 968
 <211> 313
 <212> nucleic acid
 <213> Zea mays

<400> 968

ggangtngag natnttgagg ctgntgcat tcaggcgatc cngattgatg aggcngntct 60
aagggagngn ttggcctgcg cangttatag catgcantct acctgcaactg cgctgtccnc 120
tctatcngaa tcaccnctn cagantccan nacncaacc agatccacac acatatgtgc 180
tagtnaaact ntaacnanat tatnttngga atcatcgaca tggacgcaga tgtgatcacc 240
atcgagaact cccgtccgac gagaagctac tttccgtgtt ccgngagggn gtgaagtacg 300
gagctgggat cng 313

<210> 969

<211> 222

<212> nucleic acid

<213> Zea mays

<400> 969

caaggccaaa aaagatctct gaggagggan tacgtcaactg ncatcaagga ggaaatcaag 60
gnaggtcgtt aaggggnaag aagagcttga cattgatgtg cttgtgcatg gcnagcctga 120
gagaaacgat atggttgagt acttcggtga gcaactctcg gtttttcntt nactgncaat 180
ggctnnntnc aanntantng nnnaagggnnc ntnaagnctn na 222

<210> 970

<211> 534

<212> nucleic acid

<213> Zea mays

<400> 970

agntnntnnn tggngggga aggtatgnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nncgngctcg acaccaacat nctctgggtg 120
aaccctgact gtggtctcaa gacacgcacg tncacagagg tcaagcccg ctagaccaaca 180
tggtctctgc accaagctca tncgcaccna acttgnacgc nngaaatgag gtctctngata 240
gntccatggt ctgatagcnc ggaatgagcc agntgttttg aataatttgg gtgntacccc 300
ctgnncnatg gnggaagtgn taggttaggc tctcattggn gggatacgcc gntacaagat 360
gtgttttaag nnggagtggn gagnttttct ttgngntatg nntctgggng tatgtttnac 420
gctttgntta taacagaagn tgaaatncgc anngttgcta tacttnggcc aagtnaccaa 480

agctgaccng angaanaact gcagngcacc cttacacacc aangngggaa acgg 534

<210> 971
 <211> 435
 <212> nucleic acid
 <213> Zea mays
 <400> 971

ccacgcgttc gcngacgcgt gcgggaacna ccngtctaag ttngagacat nctacnaant 60
 nncnattgca atcaaaaagg aggttgagga tcttgaggct gatggtattc aggtgatcca 120
 aatcnatngg gcanctgtaa taggagntc tggcactacn caagttacat natgcattct 180
 acctgnactg tncgtgccac tntttnaaga tcacnaactg nngatttnaa gacaccacnc 240
 agatncacac cnacatgtgc tantcnaact tcaacnacat catatactcc attatcgaca 300
 tggatgccga tntgatcacn atcgagaact cccggttntg actagaanct actgtccnga 360
 ntttcgtgan ggtatgaagt acggagctgg cattgaacct ggtgtctact acattcatct 420
 cctaggattc ccttc 435

<210> 972
 <211> 430
 <212> nucleic acid
 <213> Zea mays
 <400> 972

cccactcttc cnnccactct tncgngnttt angacaaaact tgtcatatna ncttctctggg 60
 ggctcattca caccgntgng gatcttctta acnanactaa tcttgacacc gaaattaant 120
 cttgncttgc ttttgctgcc cagaaagctg tngaggttna tgctcttgct aatgcattgg 180
 ctgntcaaaa agatnangct tacttcgcan caaatgcttc tgctcatcct cgannaaann 240
 ctcaccccggt gtgaccaatn aaaaantcca taangctnct gntgctntca aagctcttac 300
 cacctccgtg ngaccaacat taactgctag attggatgcc ccatcaaaaa naatcttaac 360
 cttcccatte cttccancac tacaatcggn tccatccaca aaccogttna acctcatnaa 420
 gatccccngt 430

<210> 973
 <211> 266
 <212> nucleic acid

<213> Zea mays

<400> 973

ttcgacttca ggccccgggat gatcagcatc aacctcgacc tgnnngaagn gcggcaacaa 60
 nttcatcaag accgccgcct aacggccact tcggccgtga cgacgccgac nttcacctgg 120
 gaggtggtga agccccctnaa gttcgacaag gcatcggntt aaggttggga gtgtcaactgt 180
 ggncatgagg actancttcc tctggctctt ctgttacctg naagattgct gctgntggat 240
 gagtgtgttt gatcatgact ggctgc 266

<210> 974

<211> 303

<212> nucleic acid

<213> Zea mays

<400> 974

gttcaactgcc aacggatggg tgcaatccta tggatcacgc tgtgtgaagc caccattat 60
 ctacggtgat gtcagccggc cgaaccccat gactgttttc tggccaaga tggcacagag 120
 catgaccct cgtcccatga agggaatgtt gactggtccg gtcacaatcc tcaactggtc 180
 attcgtcagg aacgaccagc ctaggtttga gacatgctac caaatagctc ttgcaatcaa 240
 aaaggagggtt gaggatcttg aggctgctgg tattcagggtg atccagatcg atgaggcagc 300
 tct 303

<210> 975

<211> 296

<212> nucleic acid

<213> Zea mays

<400> 975

caacggatgg gtgcaatcct atggatcacg ctgtgtgaag ccacccatta tctatggtga 60
 tgtcagccgg ccgaacccca tgactgtntt ctggccaag atggcacaga gcatgacccc 120
 tcgtcccatg aagggaatgt tgactggtcc ggtcacaatc ctgaactggt cattcgtcag 180
 gaacgaccag ctaggttttg agacatgcta ccaaataagct cttgcaatca aaaaggagggt 240
 tgaggatctt gaggctgctg gtattcagggt gatccagatc gatgaggcag ctctaa 296

<210> 976

<211> 279
<212> nucleic acid
<213> Zea mays

<400> 976

cactgccaac ggatgggtgc aatcctatgg atcacgctgt gtgaagccac ccattatcta 60
tggatgatgtc agccggccga accccatgac tgttttctgg tccaagatgg cacagagcat 120
gacccctcgt cccatgaagg gaatgttgac tggtcgggtc acaatcctga actggtcatt 180
cgtcaggaac gaccagccta ggtttgagac ntgctacaa ntagctcttg cantcaaaaa 240
ggaggttgag gntcttgagg ctgctggtat tcaggtgat 279

<210> 977
<211> 220
<212> nucleic acid
<213> Zea mays

<400> 977

ggaacggncg agnacgctg ggngtncagc cggccgaacc ccatnactgt tttctggtgc 60
aagacggcac agagcatgac cgctcgctcc atgaaggga tgttgactgg tccagtcaca 120
atcctcaact ggtcatttgt caggaacgat cagcctaggt ttgagacatg ctaccaata 180
gctcttgcaa taaaaaagga ggttgaggat cttgaggctg 220

<210> 978
<211> 295
<212> nucleic acid
<213> Zea mays

<400> 978

cctgaacttg gcccaagcac caagttcaca tacgcttctc acaaggctgt ttctgagtac 60
aaggaggcaa aggcgctcgg cattgataca gtcccagtcg ttgttgacc agtgatcac 120
ttgtctctct ctaagcctgc caagggtgtg gagaaatctt tctctctctt ttnacttctt 180
ggtagcattc ttcccatcta caaggagggt gttgctgagc tgaaggcagc tgggtgcttca 240
tggattcaac ttgatgagcc tacccttggt aaagaccttg atgctcncga attgg 295

<210> 979
<211> 259
<212> nucleic acid

<213> Zea mays

<400> 979

cttctcaciaa ggctgtttct gagtacaagg aggcaaaggc gctcggcatt gatacagtcc 60
cagtgtttgt tggaccagtg tcataacttg tcctctctaa gcctgccaaag ggtgtggaga 120
aatctttctc tcttctttca cttcttggtg gcattcttcc catctacaag gaggttgttg 180
ctgagctgaa ggcagctggt gcttcatgga ttcaacttga tgagcctacc cttgttaaag 240
accttgatgc tcacgaatt 259

<210> 980

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 980

gtttctgagt acaaggaggc aaaagcgctc ggtattgata ctgtcccagt gcttgttgga 60
ccagtgtcat acttgctcct ctctaagcct gccaaagggtg tggagaaatc tttctctctt 120
ctttcgcttc ttggcagcat tcttcccatc tacaaggagg ttgtttctga actgaaggca 180
gctgggtgctt catggattca gtttgatgag cctacccttg ttaaagacct tgatgttcac 240
gaattgggca gcattctctt cagcatatgt gaactagagt cggcatctct ggattgaatg 300
tg 302

<210> 981

<211> 284

<212> nucleic acid

<213> Zea mays

<400> 981

ggcagctggt gcttcatgga ttcagtttga tgagcctacc cttgttaaag accttgatgt 60
tcacgaattg gcagcattct cttcagcata tgctgaacta gagtcggcat tctctggatt 120
gaatgtgctt atcgagacat actttgctga cattcctgct gagtcctaca agaccctcac 180
gtcattgagt ggtgtgactg cttacgggtt tgatcttctc cgtggagcca agacccttga 240
tcttatcagg agcagttccc ctctgggaag tacctcttct ctgg 284

<210> 982

<211> 314
 <212> nucleic acid
 <213> Zea mays

 <400> 982

 tgctgagctg aangcagctg gtgcntcatg gattcaactt gatgagccta nccntgttaa 60
 agaccttgat gctcacgaat tggccgcatt ctcttcagca tatgctgaac tggagtcac 120
 gttctctgga ttgaatgtgc ttatcgagac atacttcgct gacattcctg ctgagtccta 180
 caagaccctc acatcaatga gtgggtgtga ctgcctacgg ntccgatctt aaccgtngag 240
 ccaagaccct tgatcttate aggagcagct tcccctctgg gaagtacctc ttcgtggtgt 300
 tgtagatgga cgca 314

<210> 983
 <211> 231
 <212> nucleic acid
 <213> Zea mays

 <400> 983

 atacaaacta ccactttatt gtccctgaac ttggcccaag caccaagttc acatacgtt 60
 ctcacaaggc tgtttctgag tacaaggagg caaaggcgct cggcattgat acagtcccag 120
 tgcttggttg accagtgtca tacttgctcc tctctaagcc tgccaagggt gtggagaaat 180
 ctttctctct tctttcactt cttggtagca ttcttcccat ctacaaggag g 231

<210> 984
 <211> 229
 <212> nucleic acid
 <213> Zea mays

 <400> 984

 gtttcngagt acaaggaggc aaaggcgctc ggcattgana cagtcccagt gncttgtnng 60
 accagtgtca tacttgctcc tctctaagcc tgccaagggt gtggagaaat ctttcncnct 120
 tctttcacnt cttggtagca tncntcccat ctacaaggag gttgtngcng agcngaaggc 180
 agcnggtgct tcatggattc aacttgatga gcctaccctt gttaaagac 229

<210> 985
 <211> 300
 <212> nucleic acid

<213> Zea mays

<400> 985

atctttctct ottttttcac ttcttggtag cattcttccc atctacaagg aggttggtgc 60
tgagctgaag gcagctggtg cttcatggat tcaacttgat gagcctaccc ttgttaaaga 120
ccttgatgct caccgaattg gccgcattct cttcagcata tgctgaactg gagtcacgt 180
tctctggatt gaatgtgctt atcgagacat acttcgctga cattcctgct gagtccctac 240
aagaccctca catcattgag tgggtgtgact gcttacgggt tcgatcttat ccgtggagcc 300

<210> 986

<211> 177

<212> nucleic acid

<213> Zea mays

<400> 986

gaaatctttc tctcttcttt cgcttcttgg cagcattctt cccatctaca aggagggttg 60
ttctgaactg aaggcagctg gtgcttcatg gattcagttt gatgagccta cccttgtaa 120
agaccttgat gttcacgaat tggcagcatt ctcttcagca tatgctgaac tagagtc 177

<210> 987

<211> 145

<212> nucleic acid

<213> Zea mays

<400> 987

ggaggntggt gctgagctga aggcagctgg tgcttcatgg attcagcttg atgagcctac 60
ccttggtaaa gaccttgatg ctacgaatt ggccgcattc tcttcagcat atgctgaact 120
ggagtctngt tctctggatt gaatg 145

<210> 988

<211> 189

<212> nucleic acid

<213> Zea mays

<400> 988

attcaacttg atgagcctac ccttggtaaa gaccttgatg ctacgaatt ggccgcattc 60
tcttcagcat atnctgaact ggagtcacg ttctcnggat tgaatgtgct atcgagacan 120

acttcgctga catnctgctg agtcctacaa gaccctcaca tcatgagtgg tgtgatnctt 180
acggtttctg 189

<210> 989
<211> 532
<212> nucleic acid
<213> Zea mays

<400> 989

gaggctcgnt ntnntnnngn aacttnaagg gagncttaat ggaaagcttg taccgggtccg 60
gattccccggg tcgacccacg cgtccgcgga cgcgtgggta caggcttact tcgcagcaaa 120
tgctgctgct caggcctcga ggaaatcctc accccgtgtg accaatgaag aagtccagaa 180
ggctgctgct gctctcaagg gctctgacca ccgcgtgcg accaacgtta gtgctagatt 240
ggatgccag cagaagaage ttaacctccc catccttccc accactacaa tcggttcggt 300
cccacaaacc gttgagctca ggagggtccg ccgtgagtac aaggcaaaaa agatctctga 360
ggaggaatac gtcactgcc acaaggagga aatcaacaag gtcgttaagc tccaagaaga 420
gcttgacatt gatgtgcttg tgcatggcga gcctgagaga aacgatatgg ttgagtactt 480
cggtgagcag ctctccgggtt ttgcgttcac tgccaatggc tgggtgcagt ct 532

<210> 990
<211> 441
<212> nucleic acid
<213> Zea mays

<400> 990

gcttacttcg cagcaaatgc tgctgctcag gcctcgagga aatnctnacc ccgtgtgacc 60
aatgaagaag tccagaaggc tgctgctgct ctcaagggtc ctgaccaccg ccgtgggacc 120
aacgttagtg ctagattgga tgcccagcag aagaagctta acctccccat ccttcccacc 180
actacaatcg gttcgttccc acaaaccgtt gagctcagga gggtcgcgcg tgagtacaag 240
gcaaaaaaga tctctgagga ggaatacgtc actgccatca aggaggaaat caacaaggtc 300
gttaagcttc aagaagagct tgacattgat gtgcttgtgc atggcgagcc tgagagaaac 360
gatatgggtg agtacttcgg tgagcagctc ttcggttttg cgttccactgg caatggctgg 420
gtncagtcctt atgggtcaag g 441

<210> 991
 <211> 524
 <212> nucleic acid
 <213> Zea mays

 <400> 991

 gnnnnnnnagt agtttatnnn ggggnngggg gggggaangg gaagnttcac cggtcoggaa 60
 ttcccggttc gaccacgcg tccgatccag gacaccaccc agatccacac ccacatgtgc 120
 tactccaact tcaacgacat catncactng atcatcgaca tggacgcgga cgtgatcacc 180
 atcgagaact cgcggtccga cgagaagctg ctctccgtgt tccgtgaggg tgtcaagtac 240
 ggcgcgggca tcggccccgg tgtctacgac atccactccc ccaggatccc gtcggccgag 300
 gagatcgccg accgcatcga caagatgctg gccgtgctgg acaccaacat cctctgggtg 360
 aaccccgact gcggcctcaa gaccgcgaag tacacggagg tcaagccgc cctgaccaac 420
 atggtctccg ctgctaagct cattcgcacc cagctcgcca agcgccaaag tgagcacctt 480
 tttttgctt tttcgtttcc gaggagggcg tcgtcgatgc caat 524

<210> 992
 <211> 436
 <212> nucleic acid
 <213> Zea mays

 <400> 992

 gaattnccaa ctgcgatttc caggacacca cccagatcca caccacntg tgctactcca 60
 acttcaacga catcatccac tcgatcatcg acatggacgc ggacgtgatc accattgaga 120
 actcgcggtc cgacgagaag ctgctctccg tgttccgcga ggcgtcaag tacggcgcgg 180
 gcatcgcccc cgggtgtctac gacatccact ccccaggat cccgtcggct gaggagatcg 240
 ccgaccgcat cgacaagatg ctggcgggtgc tggacaccaa catcctctgg gtgaaccccg 300
 actgcggcct taagaccgc aagtacacgg aggtcaaagc ccgcctgacc aacatggtct 360
 ncgncgctaa gctcatncgc acccagntcg ccancgcaa gtgancacct ttttttttgg 420
 cctttttngn ttccga 436

<210> 993
 <211> 330
 <212> nucleic acid

<213> Zea mays

<400> 993

gcagaaaaag ctcaaccttc ctgtccttcc cacaaccaca attggttcat tccctcagac 60
tgtggaactc aggaggggtcc gtcgtgagta caaggcaaag aagatctccg aggaggaata 120
catcagtgcc atcaaggaag aaatcagcaa ggttgtcaag atccaagagg agcttgacat 180
tgatgtgctt gtgcatggag agccagaaaag aaatgacatg gttgagtact tcggtgagca 240
attatctggt tttgcattca ctgccaacgg atgggtgcaa tcttatggat cacgttgtgt 300
gaagccaccc attatctatg gcgatgtcag 330

<210> 994

<211> 438

<212> nucleic acid

<213> Zea mays

<400> 994

atgacgtctc gcccaatgaa ggggnttctg actggcccag tcacaatcct caactgggcc 60
ttcgtccgga atgaccagcc gaggtttgag acttgctacc agatcgctct tgcgatcaag 120
aaggaagtgc aggatcttga ggctgggtgt attcaggtta tccaaatcga cgaggctgca 180
ctgagagaag ggctgccgct ccgcaaggct gaacatgcat tctacttga ctgggctgcc 240
actccttcag aattaccaac tgcgagatcc angacaccac ccagatccac acccacatgt 300
gtacttcaa cttcaacgac atcatncact cgatcatcga catggacgcc gacgtgaaca 360
ccattgagaa cttgcgggcc gacnagaaac ttntnttccg ngttncnoga aggccgtnaa 420
gtccgggncc gggcatnt 438

<210> 995

<211> 469

<212> nucleic acid

<213> Zea mays

<400> 995

cacagagcat gacctctgt cccatgaagg tnatgttgac tggctntntc acaatcctga 60
actggtcatt cgtcaggaaac gaccagccta ggtttgagac atgctaccaa atagctcttg 120
caatcaaaaa ggaggttgag gatcttgagg ctgctggat tcaggatgac cagatcgatg 180

aggcagctct aagggagggg ctgccactac gcaagtcaga gcatgcattc tacctggact 240
 gggctgtcca ctctttcagg atcaccaact gcggagtcca ggacaccacc cagatccaca 300
 cccacatgtg ctactccaac ttcaacgaca tcatccactc catcatcgac atggatgccc 360
 gatgtgatca cgatcgagaa ctcccgggtct gaccagaagc tactgtccgt cttnccgtgag 420
 ggtgtgaaat accggaactg gcattggccc tgggtgtctac gacattcat 469

<210> 996
 <211> 353
 <212> nucleic acid
 <213> Zea mays

<400> 996

tggggctgtc cactctttca ggatcaccaa ctgcggagtc caggacacca ccagatcca 60
 caccacatg tgctactcca acttcaacga catcatccac tccatcatcg acatggatgc 120
 cgatgtgatc acgatcgaga actcccggtc tgacgagaag ctactgtccg tcttccgtga 180
 ggggtgtgaag tacggagctg gcattggccc tgggtgtctac gacatccact ctccaggat 240
 tccctccacc gaggagatcg cagaccgctg cgagaagatg ctgcgccgtg tcgacaccaa 300
 catcctctgg gtgaacctga ctgtggtctc aagacacgca atacacggan gtc 353

<210> 997
 <211> 452
 <212> nucleic acid
 <213> Zea mays

<400> 997

gaaatcaaca aggtcgttaa gcttcaagaa gagcttgaca ttgatgtgct tgtgcatggc 60
 gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg ttttgcggtc 120
 actgccaatg gctgggtgca gtcttatggg tcaagggtcg tcaagcctcc gatcatctac 180
 ggtgaccgtg agccgcccc accccatgac cgttttctgg tcgaagacgg cacagagcat 240
 gacgtctcgc ccaatgaagg ggatgctgac tggcccagtc acaatcctca actggtcctt 300
 cgtccggaat gaccagccga ggtttgagac ttgctaccag atccgctctt gcgatcaaga 360
 aggaaagtcg aggatcttga agctgggtggg tattcaaggg tatccaaatc gaccagggct 420
 tgcactnaaa gaaanggctt nccgnntcgn aa 452

<210> 998
 <211> 484
 <212> nucleic acid
 <213> Zea mays

<400> 998

ttnnnnggcc cngnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn ngacatcatc 60
 cactcgatca tcgacatgga cgcggaagt atcaccatcg agaactcgcg gtccgacgag 120
 aagctgctct ccgtgttccg tgaggggtgc aagtacggcg cgggcatcgg ccccggtgtc 180
 tacgacatcc actccccag gatcccgctg gccgaggaga tcgccgaccg catcgacaag 240
 atgctggccg tgctggacac caacatctc tgggtgaacc ccgactgcgg cctcaagacc 300
 cgcaagtaca cggaggtcaa gcccgccctg accaacaatgg tctccgctgc taagtcatt 360
 cgcaccagc tcgccagcgc caagtgaaca ctttttttg cttttttcgt ttccgaagaa 420
 gggcgctcgc natgccaatt ttgtttccaa taaacaaggn tnccccccn tttncnccg 480
 ttnt 484

<210> 999
 <211> 353
 <212> nucleic acid
 <213> Zea mays

<400> 999

caaggctgag catgcattct acttggaact ggctgtccac tccttcagaa ttaccaactg 60
 cgagatccag gacaccaccc agatccacac ccacatgtgc tactccaact tcaacgacat 120
 catccactcg atcatcgaca tggacgcgga cgtgatcacc attgagaact cgcggtccga 180
 cgagaagctg ctctccgtgt tccgcgaggg cgtcaagtac ggcgcgggca tcggccccgg 240
 tgtctacgac atccactccc ccaggatccc gtcgggtgag gagatcgccg accgcatcga 300
 caagatgtgg cgggtgtggac accaacaatcc tctgggtgaa cccgattgcg gct 353

<210> 1000
 <211> 303
 <212> nucleic acid
 <213> Zea mays

<400> 1000

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 tacctggact gggctgtcca ctctttcagg atcaccaact gcgaggtcca ggacaccacc 120
 cagatccaca cccacatgtg ctactccaac ttcaacgaca tcatccactc catcatcgac 180
 atggatgccg atgtgatcac gatcgagaac tcccggctcg acgagaagct actgtccgtc 240
 ttccgtgagg gtgtgaagta cggagctggc attggccctg gtgtctacga catccactct 300
 cct 303

<210> 1001
 <211> 347
 <212> nucleic acid
 <213> Zea mays

<400> 1001

atgaggetta ctttgcagcc aatgctgctg ctcaggcctc aaggagatca tcacctagag 60
 tgacaaacga ggaggtccaa aaggctgcag ctgctttgaa gggttctgac ncccgccgtg 120
 ctactactgt tgctgctaga ttggatgctc agcagaaaaa gctcaacctt cctgtccttc 180
 ccacaaccac aattgnntca ttccctcaga ctgtggaact caggagggtc cgctcgtgagt 240
 acaaggcaaa gaagatctcc gaggaggaat acatcagtgc catcaaggaa gaaatcagca 300
 aggttgtcaa gatccaagag gagcttgaca ttgatgtgct tgtgcat 347

<210> 1002
 <211> 319
 <212> nucleic acid
 <213> Zea mays

<400> 1002

caaggcaaag aagatctccg aggaggaata catcagtgcc atcaaggaag aaatcagcaa 60
 ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag agccagaaag 120
 aaatgacatg gttgagtact tcggtgagca attatctggt tttgcattca ctgccaacgg 180
 atgggtgcaa tottatggat cacgttgtgt gaagccaccc attatctatg gcgatgtcag 240
 cggccgaac cccatgactg ttttctggtc caagacggca cagagcatga ccgctcgtcc 300
 catgaaggga atgttgact 319

<210> 1003

<211> 379
 <212> nucleic acid
 <213> Zea mays

 <400> 1003

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 gactaagctt gacagcgaga ttaagtcttg gcttgctttt gctgcccaga aggttggtga 120
 gggtgatgct cttgctaagg cattggctgg tcaaaaggat gaggttact tcgcagcaaa 180
 tgctgctgct caggcctcga ggaaatcctc accccgtgtg accaatgaag aagtccagaa 240
 ggctgctgct gctctcaagg gctctgacca ccgccgtggg accaacgtta gtgctagatt 300
 ggatgccag cagaagaagc ttaacctccc catccttccc accactacaa tcgggttcgg 360
 nnnnnnnnnn nnnnnnnnt 379

1004
 351
 nucleic acid
 Zea mays
 1004

<210> 1004
 <211> 351
 <212> nucleic acid
 <213> Zea mays

 <400> 1004

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 tccatcatcg acatggatgc cgatgtgatc acgatcgaga actcccggtc tgacgagaag 120
 ctactgtccg tcttccgtga ggggtgtgaag tacggagctg gcattggccc tgggtgtctac 180
 gacatccact ctctaggat tccctccacc gaggagatcg cagaccgcgt cgagaagatg 240
 ctgccgtgc tcgacaccaa catcctctgg gtgaaccctg actgtggtct caagacacgc 300
 aatacacgga ggtcaagccc gncctgacca acagggtctcg gcaacaagct c 351

<210> 1005
 <211> 302
 <212> nucleic acid
 <213> Zea mays

 <400> 1005

 atcgacgagg ctgcaactgag agaagggctg ccgctccgca aggctgagca tgcattctac 60
 ttggactggg ctgtccactc cttcagaatc accaactgcg agatccagga caccacccag 120
 atccacaccc acatgtgcta ctccaacttc aacgacatca tccactcgat catcgacatg 180

gaagcgggacg tgatcaccat cgagaactcg cgggccgacg agaagctgct ctccgtgttc 240
 cgtgaggggtg tcaagtacgg cgcgggcatc ggccccgggtg tctacgacat ccactccccc 300
 ag 302

<210> 1006
 <211> 296
 <212> nucleic acid
 <213> Zea mays
 <400> 1006

gcagaaaaag ctcaaccttc ctgtccttcc cacaaccaca attggttcat tccctcagac 60
 tgtggaactc aggaggggtcc gtcgtgagta caaggcaaag aagatctccg aggaggaata 120
 catcagtgcc atcaaggaag aaatcagcaa ggttgtcaag atccaagagg agcttgacat 180
 tgatgtgctt gtgcatggag agccagaaaag aaatgacatg gttgagtact tcggtgagca 240
 attatctggt tttgcattca ctgccaacgg atgggtgcaa tcttatggat cacgtt 296

<210> 1007
 <211> 537
 <212> nucleic acid
 <213> Zea mays
 <400> 1007

gntngagtcn ggcttcggcn ttgatgatng cgtantcntt cngnangccg ntngaattc 60
 ccgggtcgac ccacgcgtcc ggnaggaata cgtcactgcc atcaaggagg aaattaacaa 120
 ggtcgttaag cttcaagaag agcttgacat tgatgtgctt gtgcatggng agcctgagag 180
 aaacgatatg gttgagtact tcggtgagca gctctncggt tttgcgttca ctgccaatgg 240
 ctgggtgcag tcttatgggt caaggtgcgt caagcctccg atcatctacg gtgacgtgag 300
 ccgtcccaac cccatgaccg ttttctggtc gaagacggcn canagcatga cgtctcgccc 360
 aatgaaaggg gatgctgact ngcccagtca caatcctcaa ctggtncctt gtcccggaat 420
 naccagccca gggtttgaga nttgctacca gaatcggctt cttgcgatca agaaaggga 480
 gtncaaggga tctttgangn tgggtggtnt ttaangttta ttncaaaatc gacnaag 537

<210> 1008
 <211> 347
 <212> nucleic acid

<213> Zea mays

<400> 1008

cgttttctgg tcgaagacgg ggcagagca tgacgtctcg cccaatgaag gggatgctga 60
ctggcccagt cacaatcctc aactggtcct tcgtccggaa cgaccagccg aggtttgaga 120
cttgctacca gatcgctctt gcgatcaaga aggaagtcga ggatcttgag gctgggtgga 180
ttcaggttat ccaaatcgac gaggtgcac tgagagaagg gctgccgctc cgcaaggctg 240
agcatgcatt ctacttgga cttgggtgtcc actccttcag aattaccaac tgcgagatnc 300
aggacaccac cagatccaca cccacatgtg ctactccaac ttcaacg 347

<210> 1009

<211> 339

<212> nucleic acid

<213> Zea mays

<400> 1009

gaggatcttg aggctgctgg tattcagggtg atccagattg atgaggcagc tctaagggag 60
ggtctgccac tgcgcaagtc agagcatgca ttctacctgg actgggctgt ccaactcttc 120
agaatcacca actgcgaggt ccaggacacc acccagatcc acacccatat gtgctactcc 180
aacttcaacg acatcatcca ctccatcatc gacatggacg cagatgtgat caccatcgag 240
aactcccggt ccgacgagaa gctactgtcc gtgttccgag aggtgtgaag tacggagctg 300
gaatcgggtcc agtgtctagg aatccactcg ccaagtccc 339

<210> 1010

<211> 313

<212> nucleic acid

<213> Zea mays

<400> 1010

ccgctccgca aggctgagca tgcattctac ttggactggg ctgtccactc cttcagaaac 60
accaactgcg agatccagga caccacccag atccacaccc acatgtgcta ctccaacttc 120
aacgacatca tccaatcgat catcgacatg gacgaggacg tgatcaccat cgagaactcg 180
cggctccgacg agaagctgct ctccgtgttc cgtgagggtg tcaagtacgg cgcgggcatc 240
ggccccggtg tctacgacat ccaactcccc aggatcccggt cggccganga gatcgccgac 300

cgcacgcaca aga

313

<210> 1011
<211> 310
<212> nucleic acid
<213> Zea mays

<400> 1011

caacaaggtc gttaagcttc aagaagagct tgacattgat gtgcttgtgc atggcgagcc 60
tgagagaaaac gatatggttg agtacttcgg tgagcagctc tccggttttg cgttcactgc 120
caatggctgg gtgcagtctt atgggtcaag gtgcgtcaag cctccgatca tctacggtga 180
cgtgagccgc cccaacccca tgaccgtttt ctggtcgaag acggcacaga gcatgacgtc 240
tcgccccaatg aaggggatgc tgactggccc agtcacaatc ctcaactggt ccttcgtccg 300
gaatgaccag 310

<210> 1012
<211> 319
<212> nucleic acid
<213> Zea mays

<400> 1012

acgatatggt tgagtacttc ggtgagcagc tctccggttt tgcgttcact gccaatggct 60
gggtgcagtc ttatgggtca aggtgcgtca agcctccgat catctacggt gacgtgagcc 120
gccccaaacc catgaccgtt ttctggtcga agacggcgca gagcatgacg tctcgcccaa 180
tgaaggggat gctgactggc ccagtcacaa tctcaactg gtccttcgtc cggaatgacc 240
agccgaggtt tgagacttgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc 300
tcgaggctgg tggatttca 319

<210> 1013
<211> 315
<212> nucleic acid
<213> Zea mays

<400> 1013

ctagattgga tgcccagcag aagaagctta acctcccat ccttcccacc actacaatcg 60
gttcgttccc acaaaccgtt gagctcagga gggtcgccg tgagtacaag gcaaaaaaga 120

tctctgagga ggaatacgtc actgccatca aggaggaaat caacaaggtc gttaagcttc 180
aagaagagct tgacattgat gtgcttgtgc atggcgagcc tgagagaaac gatatggttg 240
agtacttcgg tgagcagctc tccggttttg cgttcaactgc caatggctgg gtgcagtctt 300
atgggtcaag gtgcg 315

<210> 1014
<211> 505
<212> nucleic acid
<213> Zea mays
<400> 1014

aggggtgtnag gggggggaag aaaggaactg tcgtaccggt ccggaattcc cgggtcgacc 60
cacgcgtccg cttgtccgtg gaacccaaac tcttgggctt gtcacgagtg ctggtttccc 120
tgctggaaaag tacctctttg ctgggtgttg ggatggacgc aacatctggg ctgatgatct 180
tgctacatct ctcagcactc tccagtctct tgaggctgtt gttgggaagg acaagcttgt 240
cgtatcaact tctgtctgc tcatgcacac cgctgtggat cttgtgaacg agactaagct 300
tgacagcgag attaatctt ggcttgctt tgctgccag aaggttgttg aggttgatgc 360
tcttgctaag gcattggctg gtcaaaagga tgaggcttac ttcgcagcaa atgctgctgc 420
tcaagcctcg aggaaatcct naccctgtgt gaccaatgaa aaatccagaa ngctgctgct 480
gntcttinaag gcttttgacc accgc 505

<210> 1015
<211> 298
<212> nucleic acid
<213> Zea mays
<400> 1015

gccgtgaata caaggcaaag aagatcaccg aggacgaata catcagtgcc atcaaggaag 60
aaatcagcaa ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag 120
agccagagag aaatgacatg gttgagtact tcggtgagca attatctggt ttgctgttca 180
ctgccaacgg atgggtgcaa tcctatggat cacgctgtgt gaagccaccc attatctatg 240
gtgatgtcag ccggccgaac cccatgactg ttttctggtc caagatggca cagagcat 298

<210> 1016

<211> 493
 <212> nucleic acid
 <213> Zea mays
 <400> 1016
 gggnnnnnngn aangttacgt tncgggnccg gtccaaaata aacgggtcga ngcacgcgtc 60
 cgggccaaact tgtcctccaa ccttgattcc tgacaaaatt ggctggcatt ctctgctgca 120
 tacgcagaaa cttgaatctg tactttctgg attgaatgtg cttgttgaga cttactttgc 180
 tgatgttcct gctgagtcct acaagaccct aacatctctg agcagtgatga ctgcttatgg 240
 ttttgatctt gtccgtggaa cccaaactct tgggcttgtc acgagtgtg gtttccctgc 300
 tggaaagtac ctctttgctg gtgttggtga tggacgcaac atctgggctg atgatcttgc 360
 tacatctctc agcactctcc agtctcttga ggctgttggt gggaaggaca agcttgctgt 420
 atcaacttcc tgctcgctca tgcacaccgc tgtggatctt gtgaacgaga ctaagcttga 480
 cagcgagatt aag 493

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<210> 1017
 <211> 356
 <212> nucleic acid
 <213> Zea mays
 <400> 1017
 cgccgctgcg ccgtttctct ccgctcccgt cgtccggaaa gatggcgtcc cacattgttg 60
 gataccctcg catgggcccc aagagggagc tcaagtttgc acttgagtct ttctgggatg 120
 ggaagagcac cgctgaggat ctggagaagg ttgctaccga cctcagggcc agtatctgga 180
 agcagatggc tgatgctggg atcaagtaca tccccagcaa cactttctcg tactatgacc 240
 aggtccttga caccactgcc atgctcggcg ctgtccccga acgttactca tggactggan 300
 gagagattgg anttgacacc tactttctcca tggccagggc aacgcactgt tcctgc 356

<210> 1018
 <211> 344
 <212> nucleic acid
 <213> Zea mays
 <400> 1018
 gcctaccatc tggtcagaca agcttgctgt atcaacttcc tgctcgctca tgcacaccgc 60

tgtggatctt gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgcttttgc 120
 tgcccagaag gttgttgagg ttgatgctct tgctaaggca ttggctgggc aaaaggatga 180
 ggcttacttc gcagcaaag ctgcagctca ggctcgagg aaatcctcac cccgtgtgac 240
 caatgaagaa gtccagaagg ctgctgctgc tctcaagggc tctgaccacc gccgtgagac 300
 caacgttagt gctagattng atgccagca gaagaagcct aacc 344

<210> 1019
 <211> 292
 <212> nucleic acid
 <213> Zea mays

<400> 1019

ccgccgtgct accactgttg ctgctagatt ggatgctcag cagaaaaagc tcaaccttcc 60
 tgtccttccc acaaccacaa ttggttcatt cctcagact gtggaactca ggagggtccg 120
 tcgtgagtac aaggcaaaga agatctccga ggaggaatac atcagtgccca tcaaggaaga 180
 aatcagcaag gttgtcaaga tccaagagga gcttgacatt gatgtgcttg tgcattggaga 240
 gccagaaaga aatgacatgg ttgagtactt cggtgagcaa ttatctggtt tt 292

<210> 1020
 <211> 312
 <212> nucleic acid
 <213> Zea mays

<400> 1020

accagccgag gtttgagact tgctaccaga tcgctcttgc gatcaagaag gaagtcgagg 60
 atctcgaggc tgggtgtatt caggttatcc aaatcgacga ggctgcactg agagaagggc 120
 tgccgctccg caaggctgag catgcattct acttgactg ggctgtccac tccttcagaa 180
 tcaccaactg cgagatccag gacaccaccc agatccacac ccacatgtgc tactccaact 240
 tcaacgacat catccactcg atcatcgaca tggacgcgga cgtgatcacc atcgagaact 300
 cgcggtccga cg 312

<210> 1021
 <211> 286
 <212> nucleic acid
 <213> Zea mays

<400> 1021

caagatccaa gaggagcttg acattgatgt gcttgtgcat ggagagccag agagaaatga 60
catggttgag tacttcggtg agcaattatc tggttttgcg ttcaactgcc aaggatgggt 120
gcaatcctat ggatcacgct gtgtgaagcc acccattatc tacggtgatg tcagccggcc 180
gaaccccatg actgttttct ggtccaagat ggcacagagc atgacccctc gtcccatgaa 240
gggaatggtg actgggtccg tcacaatcct caactggtca ttcgtc 286

<210> 1022

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1022

atgtgctact ccaacttcaa cgacatcatc cactccatca tcgacatgga tgccgatgtg 60
atcacgatcg agaactcccg atctgacgag aagctactgt ccgtcttccg tgaggggtgtg 120
aagtacggag ctggcattgg ccttggtgtc tacgacatcc actctcctag gattccctcc 180
accgaggaga tcgcagaccg cgtcgagaag atgctcgccg tgctggacac caacatcctc 240
tgggtgaacc ctgactgtgg tctcaagaca cgcaagtaca cggaggt 287

<210> 1023

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 1023

caaggntgtc aagatccaag aggagcttga cattgatgtg cttgtgcatg gagagccaga 60
gagaaatgac atggttgagt acttcggtga gnaattatct ggttttgagt tcaactgcca 120
cggatgggtg caatcctatg gatcacgctg tgtgaagcca cccattatct atggtgatgt 180
cagccggccg aaccccatga ctgttttctg gtccaagatg gcacagagca tgacccctcg 240
tcccatgaag ggaatgttga ctgggtccggt cacaatcctg aactggtcat tcgtc 295

<210> 1024

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1024

agaagatcac cgatgacgaa tacatcagtg ccatcaagga agaaatcagc aaggttgtca 60
agatccaaga ggagcttgac attgatgtgc ttgtgcatgg agagccagag agaaatgaca 120
tggttgagta cttcgggtgag caattatctg gttttgcggt cactgccaac ggatgggtgc 180
aatcctatgg atcacgctgt gtgaagccac ccattatcta tggatgatgtc agccggccga 240
accccatgac tgttttctgg tccaagatgg cacagagcat gacccctcgt cccatgaag 299

<210> 1025

<211> 312

<212> nucleic acid

<213> Zea mays

<400> 1025

tgagtacaag gcaaaaaaga tctctgagga ggaatacgtc actgccatca aggaggaaat 60
caacaaggtc gttaagcttc aagaagagct tgacattgat gtgcttgtgc atggcgagcc 120
tgagagaaac gatatggttg agtacttcgg tgagcagctc tccggttttg cgttcactgc 180
caatggctgg gtgcagtctt atgggtcaag gtgcgtcaag cctccgatca tctacggtga 240
cgtgagccgc cccaacccca tgaccgtttt ctggtcgaag aggcacagag catgacgtct 300
cgcccaatga ag 312

<210> 1026

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1026

aaagatctct gaggaggaat acgtcactgc catcaaggag gaaatcaaca aggtcgttaa 60
gcttcaagaa gagcttgaca ttgatgtgct tgtgcatggc gagcctgaga gaaacgatat 120
ggttgagtac ttcggtgagc agctctccgg ttttgcgttc actgccaatg gctgggtgca 180
gtcttatggg tcaaggtgcg tcaagcctcc gatcatctac ggtgacgtga gccgccccaa 240
cccatgacc gttttctggt cgaagacggc acagagcatg acgtctcgcc caatgaagg 299

<210> 1027

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 1027

gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggctct gaccaccgcc 60
gtgggaccaa cgttagtgt agattggatg cccagcagaa gaagcttaac ctccccatcc 120
ttcccaccac tacaatcggg tegtccccc aaaccgttga gtcaggagg gtccgcccgtg 180
agtacaaggc aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca 240
acaaggctgt taagcttcaa gaagagcttg acattgatgt gcttgtgcat ggcgagcctg 300
agagaaacgg 310

<210> 1028

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 1028

gtctcgccca atgaagggga tgctgactgg ccagtcaca atcctcaact ggtccttcgt 60
ccggaatgac cagccgaggt ttgagacttg ctaccagatc gctcttgcca tcaagaagga 120
agtgcaggat cttgaggctg gtggtattca ggttatccaa atcgacgagg ctgcactgag 180
agaagggctg ccgctccgca aggctgagca tgcattctac ttggactggg ctgtccactc 240
cttcagaatt accaactgcg agatccagga caccaccag atccacacc acatgtgcta 300
ctcca 305

<210> 1029

<211> 330

<212> nucleic acid

<213> Zea mays

<400> 1029

cgctgaggat ctggagaagg ttgctaccga cctcagggcc agtatctgga agcagatggc 60
tgatgctggg atcaagtaca tccccagcaa caccttctcg tactatgacc agntccttga 120
caccactgcc atgctcggcg ctgtccccga acgctactca tggactggag gagagattgg 180
gtttgacacc tactttctcca tggccagggg caacgccact gtccctgcta tggagannac 240
caagtggttt gacaccaact accactttat tgtccctgaa ttgggcccaa acaccaagtt 300

ctcctatgct tctcacaagg ctgttaacga

330

<210> 1030
<211> 269
<212> nucleic acid
<213> Zea mays

<400> 1030

gcattctacc tggactgggc tgtccactct ttcagaatca ccaactgagg agtccaggac 60
accacccaga tccacaccca tatgtgtctac tccaacttca acgacatcat ccactccatc 120
atcgacatgg acgcagatgt gatcaccatc gagaactccc ggtccgacga gaagctactg 180
tccgtgttcc gcgaggggtgt gaagtacgga gctggaatcg gtccaggtgt ctacgacatc 240
cactcgccca ggatcccctc caccgagga 269

<210> 1031
<211> 311
<212> nucleic acid
<213> Zea mays

<400> 1031

ctcgcccaat gaaggggatg ctgactggcc cagtcacaat cctcaactgg tccttcgtcc 60
ggaatgacca gccgaggttt gagacttgct accagatcgt cttgcgatca agaaggaagt 120
cgaggatctc gaggctggtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 180
aggggtgccg ctccgcaagg ctgagcatgc attctacttg gactgggctg tccactcctt 240
cagaatcacc aactgcgaga tccaggacac caccagatc cacaccaca tgtgtctactc 300
caacttcaac g 311

<210> 1032
<211> 382
<212> nucleic acid
<213> Zea mays

<400> 1032

gggacaccac ccagatccac acccacatgt gctactccaa ctttcaacga catcatccac 60
tcgatcatcg acatggacgc ggacgtgatc accattgaga actcgcggtc cgacgagaag 120
ctgctctccg tgntccgcga gggcgtcaag tacggcgcgg gcatcggcnc cggtgtctac 180

gacatccact cccccaggat cncgtcggct gangagatcg ccgaccgcat cgacaagatg 240
 ctggcgggtgc tggacancaa catcctctgg gtgaaccccg actgcngcct caagacccgc 300
 aagtacacgg agtcaagccc gentgancaa catgggtctcc gccgtaagct catccgnacc 360
 agctcgccag cgcaatgana ac 382

<210> 1033
 <211> 292
 <212> nucleic acid
 <213> Zea mays
 <400> 1033

aggatcttga ggctgggtgt attcaggtta tccaaatcga cgaggctgca ctgagagaag 60
 ggctgccgct ccgcaaggct gagcatgcat tctacttga ctgggctgtc cactccttca 120
 gaattaccaa ctgcgagatc caggacacca ccagatcca gaccacatg tgctactcca 180
 acttcaacga catcatccac tcgatcatcg acatggacgc ggacgtgatc accattgaga 240
 actcgoggto cgacgagaag ctgctctccg tgttccgcga gggcgtcaag ta 292

<210> 1034
 <211> 320
 <212> nucleic acid
 <213> Zea mays
 <400> 1034

cttgacagcg agattaagtc ttggcttgct tttgctgccc agaaggttgt tgaggttgat 60
 gctcttgcta aggcattggc tgggtcaaaag gatgaggctt acttcgcagc aaatgctgca 120
 gctcaggcct cgaggaaatc ctcaccccggt gtgaccaatg aagaagtcca gaaggctgct 180
 gctgctctca agggctctga ccaccgccgt gcgaccaacg ttagtgctag attggatgcc 240
 cagcagaaga agcttaacct cccatcctt cccaccacta caatcggttc gttcccacaa 300
 accgttgagc tcaggagggt 320

<210> 1035
 <211> 305
 <212> nucleic acid
 <213> Zea mays
 <400> 1035

acttcggtga gcagctctcc ggttttgcgt tcaactgccaa tggctgggtg cagtcttatg 60
 ggtcaagggtg cgtcaagcct ccgatcatct acggtgacgt gagccgcccc aaccccatga 120
 cegttttctg gtcgaagacg gcacagagca tgacgtctcg cccaatgaag gggatgctga 180
 ctggcccagt cacaatcctc aactggtcct tcgtccggaa tgaccagccg aggtttgaga 240
 cttgctacca gatcgctctt gcgatcaaga aggaagtcga ggatcttgag gctgggtgga 300
 ttcag 305

<210> 1036
 <211> 278
 <212> nucleic acid
 <213> Zea mays

<400> 1036
 caagaggagc ttgacattga tgtgcttggt catggagagc caganagaaa tgacatgggt 60
 gagtacttcg gtgagcaatt atctggtttt gcgttcaactg ccaacggatg ggtgcaatcc 120
 tatggatcac gctgtgtgaa gccacccatt atctatgggt atgtcagccg gccgaacccc 180
 atgactgttt tctgggtccaa gatggcacag agcatgaccc ctcggtcccat gaagggaatg 240
 ttgactggtc cggtcacaat cctgaactgg tcattcgt 278

<210> 1037
 <211> 298
 <212> nucleic acid
 <213> Zea mays

<400> 1037
 ctctgaggag gaatacgtca ctgccatcaa ggaggaaatc aacaaggtcg ttaagcttca 60
 agaagagcgtt gacattgatg tgcttggtgca tggcgagcct gagagaaacg atatggttga 120
 gtacttcggt gagcagctct ccggttttgc gttcactgcc aatggctggg tgcagtctta 180
 tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac gtgagccgcc ccaaccccat 240
 gaccgttttc tggtcgaaga cggcacagag catgacgtct cgcccaatga aggggatg 298

<210> 1038
 <211> 298
 <212> nucleic acid
 <213> Zea mays

<400> 1038

ctctgaggag gaatacgtca ctgccatcaa ggaggaaatc aacaaggctcg ttaagcttca 60
agaagagctt gacattgatg tgcttgtgca tggcgagcct gagagaaaacg atatggttga 120
gtacttcggt gaggagctct ccggttttgc gttcaactgcc aatggctggg tgcagtctta 180
tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac gtgagccgcc ccaaccccat 240
gaccgttttc tggtcgaaga cggcacagag catgacgtct cgcccaatga aggggatg 298

<210> 1039

<211> 447

<212> nucleic acid

<213> Zea mays

<400> 1039

ggcgctcggg gttgataccg tgccagtact tgttggacca gtctcgtacc tgttgcngnn 60
aaagcctgcc aaggggtgtg agaagggatt cctcttctt tcccttctta gcagcatcct 120
cccagtctac aaggagggtca ttgctgagtt gaaggcagct ggggcttcat ggattcagtt 180
tgatgagccc actcttgtcc tcgacctga ttctgacaaa ttggtgcat tctctgctgc 240
atatgcagaa cttgaatctg tactttctgg attgaatgtg cttgttgaga cttactttgc 300
tgatgttctt gctgagtcct acaagacct aacatccctg aacagtgtga ctgcttatgg 360
gtttgatent gtccggggac ccnaaactnt tgggcttggc accaaggctg gnttccttnt 420
tggaanaaac cctntttgct ggggggtg 447

<210> 1040

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 1040

ctccaagaag agcttgacat tgatgtgctt gtgcatggcg agcctgagag aaacgatatg 60
gttgagtact tcggtgagca gctcctccgg ttttgcgttc actgccaatg gctgggtgca 120
gtcttatggg tcaaggtgag tcaagcctcc gatcatctac ggtgacgtga gccgccccaa 180
ccccatgacc gttttctggt cgaagacggc gcagagcatg acgtctcgcc caatgaaggg 240
gatgctgact ggcccagtca caatcctcaa ctggctcttc gtccggaatg accagccgag 300

<210> 1041
 <211> 297
 <212> nucleic acid
 <213> Zea mays

<400> 1041

caaggtcggtt aagctccaag aagagcttga cattgatgtg cttgtgcatg gcgagcctga 60
 gagaaacgat atggttgagt acttcggtga gcagctctcc ggttttgcgt tcaactgccaa 120
 tggctgggtg cagtcttatg ggtcaagggtg cgtcaagcct ccgatcatct acggtgacgt 180
 gagccgcccc aaccccatga ccgttttctg gtcgaagagg cgcagagcat gacgtctcgc 240
 ccaatgaagg ggatgctgac tggcccagtc acaatcctca actggtcctt cgtccgg 297

<210> 1042
 <211> 548
 <212> nucleic acid
 <213> Zea mays

<400> 1042

gnnnaggtgn tnannggggn nggaaggaac nttttaccgg tccggaattc ccgggtcgac 60
 ccacgcgtcc gccacgcgt ccgctcgatc atcgacatgg acgcggacgt gatcaccatt 120
 gagaactcgc ggnccgacga gaagctgctc tccgtgttcc gcgagggcgt caagtacggc 180
 gcgggcatcg gccccggtgt ctacgacatc cactccccca ggatcccgtc ggctgaggag 240
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 cccgactgcg gcctcaagac ccgcaagtac acggaggta agcccgctg accaacaatgg 360
 tctccgccgc taagctcatc cgcacccagc ttgncagcgc caagtgagca ccnttttttt 420
 tgcccttttc ggtttcgaag aaggcgtcgt cgatgccaat ttggtttcca ataaaaaggc 480
 tttgcccgcc gtctgtgnac tccgctgngg taaggtaaga agttttctta aaaaaaaaaa 540
 angggggc 548

<210> 1043
 <211> 303
 <212> nucleic acid
 <213> Zea mays

<400> 1043

gacggcgag agcatgacgt ctgcccgaat gaaggggatg ctgactggcc cagtcacaat 60
cctcaactgg tccttcgtcc ggaatgacca gccgaggttt gagacttgct accagatcgc 120
tcttgcgatc aagaaggaag tcgaggatct cgaggctggt ggtattcagg ttatccaaat 180
cgacgaggct gcactgagag aagggctgcc gctccgcaag gctgagcatg cattctactt 240
ggactgggct gtccactcct tcagaatcac caactgcgag atccaggaca ccaccagat 300
cca 303

<210> 1044

<211> 602

<212> nucleic acid

<213> Zea mays

<400> 1044

gcggacgtga tcaccattga gaactcgcgg tccgacgaga agctgctctc cgtgttcgcg 60
gagggcgctca agtacggcgc gggcatcggc cccggtgtct acgacatcca ctccccaggg 120
atcccgtcgg ctgaggagat cgccgaccgc atcgacaaga tgctggcggt gctggacacc 180
aacatcctct ggggtgaacc cgactgcggc ctcaagacc gcaagtacac ggaggtcaag 240
cccgccctga ccaacatggt ctccgccgct aagctcatcc gcaccagct cgccagcgcc 300
aagtgagcac cttttttttg cttttttcgt ttccgaggag ggcgtcgtcg atgccaattt 360
gtttccaata aacagggctg tgcccgccgt tctgttgtag tccgtctgtg gttagggttag 420
tagttttctt gatctcgccc ccacgcggta cctgttttac tactctctgt ttggtggttt 480
ctgaggcaag ttgcccggtg acttgatcgc taaaaaccga gtgggacatc tgcacggttt 540
catctgncac cactnccccg aggtgtgctg gttcacggct caaagaaaga tgtcntatcg 600
gc 602

<210> 1045

<211> 273

<212> nucleic acid

<213> Zea mays

<400> 1045

ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag agccagagag 60

aaatgacatg gttgagtact tcggtgagca attatctggt tttgcgttca ctgccaacgg 120
 atgggtgcaa tcctatggat cacgctgtgt gaagccaccc attatctatg gtgatgtcag 180
 ccggccgaac cccatgactg ttttctggtc caagatggca cagagcatga cccctcgtcc 240
 catgaaggga atgttgactg gtccggtcac aat 273

<210> 1046
 <211> 328
 <212> nucleic acid
 <213> Zea mays

<400> 1046

atcatcgccg ctgcgcgctc tcctccgctc ccgtcgtccg gaaagatggc gtcccacatt 60
 gttggatacc ctgcgatggg ccccaagagg gagctcaagt ttgcacttga gtctttctgg 120
 gatgggaaga gcaccgctga ggatctggag aagggttgcta ccgacctcag ggccagtatc 180
 tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt ctcgactat 240
 gaccaggctc ttgacaccac tgccatgctc ggcgctgtcc ccgaacgtta ctcatgnact 300
 ggaggagaga ttggatttga cacctact 328

<210> 1047
 <211> 297
 <212> nucleic acid
 <213> Zea mays

<400> 1047

aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca acaaggctgt 60
 taagcttcaa gaagagcttg acattgatgt gcttgtgcat ggcgagcctg agagaaacga 120
 tatggttgag tacttcggtg agcagctctc cggtttggcg ttactgcca atggctgggt 180
 gcagtcttat gggtaaggt gcgtaagcc tccgatcatc tacggtgacg tgagccgccc 240
 caaccccatg accgttttct ggtcgaagac ggcacagagc atgacgtctc gcccaat 297

<210> 1048
 <211> 264
 <212> nucleic acid
 <213> Zea mays

<400> 1048

ctcttttcaga atcaccaact ggggagtcca ggacaccacc cagatccaca cccatatgtg 60
ctactccaac ttcaacgaca tcatccactc catcatcgac atggacgcag atgtgatcac 120
catcgagaac tcccgggtccg acgagaagct actgtccgtg ttccgcgagg gtgtgaagta 180
cggagctgga atcgggtccag gtgtctacga catccactcg cccaggatcc cctccaccga 240
ggagatcgcg gaccgcgtca acaa 264

<210> 1049
<211> 471
<212> nucleic acid
<213> Zea mays

<400> 1049

gggnnnnacg ttantcngcg ngnnccgtca naggacncac nggccgaggc ncgcgtccac 60
accnatcatc gccgctgcgc cgtctcctcc gctcccgtcg tccggaaaga tggcgtccca 120
cattgttgga taccctcgca tgggccccaa gagggagctc aagtttgac ttgagtcttt 180
ctgggatggg aagagcaccg ctgaggatct ggagaagggt gctaccgacc tcagggccag 240
tatctggaag cagatggctg atgctgggat caagtacatc cccagcaaca ccttctcgta 300
ctatgaccag gtcccttgaca ccactgccat gctcggcgct gtccccgaac gctactcatg 360
gactggagga gagattgggt ttgacaccta acttctccat ggccaagggc aacgccactg 420
tccctgctat gggagatacc aanttgggtt tggacacaaa ctaccaactt t 471

<210> 1050
<211> 291
<212> nucleic acid
<213> Zea mays

<400> 1050

cgctcccgtc gtccggaaag atggcgctccc acattgttgg ataccctcgc atgggccccca 60
agaggagct caagtttgca cttgagtctt tctgggatgg gaagagcacc gctgaggatc 120
tggaagaagg tgctaccgac ctgaggcca gtatctggaa gcagatggct gatgctggga 180
tcaagtacat cccagcaac actttctcgt actatgacca ggtccttgac accactgcca 240
tgctcggcgc tgtcccgaa cgttactcat ggactggagg agagattgga t 291

<210> 1051

<211> 349
 <212> nucleic acid
 <213> Zea mays

 <400> 1051

 atccactcca tcatcgacat ggatgccgat gtgatcacga tcgagaactc ccgatctgac 60
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 tacttggact gggctgtcca ctcttcaga atcaccaact gcgagatcca ggacaccacc 180
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 gtgtctacga catccactcc cccaggatcc cgtcggccga ggagatcgcc gaccgcatcg 240
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<400> 1070

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 tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120
 gctctccggt tttgcgttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180
 caagcctccg atcatctacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240
 gaagaggcac agagcatgac gtctcgccca atgaagggga tgctgactg 289

<210> 1071
 <211> 486
 <212> nucleic acid
 <213> Zea mays

<400> 1071

gctcagcaga aaaagctcaa ccttcngtc cnnccacaa ccacaattgg ttcattcccg 60

gaaacggtgg aactcaggng ggttcgncgt gaatacaagg caaagaagat caccgaggac 120
 gaatacatca gtgccatcaa ggaagaaatc agcaaggtn gcaagatcca ataggagctn 180
 gacattgatg tgcttgtgca tggagagcca gagagaaatg acatgggtga gtacttcggn 240
 gagcaattat ctgggttttag cgttcactgc caacggatgg gtgcaatnct attgatcacg 300
 ctgtggtgaa gccacccatt atctatgnga tgtcaaagcg gcccnacccc atnctgntnt 360
 ctggtcccaa gatngancat acatgacccc tttnccatg naagggaatg gttgactggg 420
 tcnggtcaca attcttaaac ttggtcattn gntaaaggaa cgacnancct taggtttnaa 480
 anannc 486

<210> 1072
 <211> 286
 <212> nucleic acid
 <213> Zea mays

<400> 1072
 aaccccatga ccgttttctg gtcgaagagg tcacagagca tgacgtctcg cccaatgaag 60
 gggatgctga ctggcccagt cacaatcttc aactggctct tcgtccggaa tgaccagccg 120
 aggtttgaga cttgctacca gntcgctctt gcgatcaaga aggaagtcga ggatcttgag 180
 gctggtggta ttacaggttat ccaaatcgac gaggtgcac tgagagaagg gctgccgctc 240
 cgcaangctg agcatgcatt ctacttgga tgggctgtcc actcct 286

<210> 1073
 <211> 305
 <212> nucleic acid
 <213> Zea mays

<400> 1073
 gaagagcttg acattgatgt gcttgtgcat ggcgagcctg agagaaacga tatggttgag 60
 tacttcggtg agcagctctc cggttttgctg ttactgcca atggctgggt gcagtcttat 120
 gggatcaagg ggcgcaagcc tccgatcatc tacggtgacg tgagccgccc caaccccatg 180
 accgttttct ggtcgaagac ggcacagagc atgacgtctc gcccaatgaa ggggatgctg 240
 actggcccag tcacaatcct caactggtct tcgtccggaa tgaccagccg aggtttgaga 300
 ttgct 305

<210> 1074
 <211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 1074

 caatcgggttc gttcccacaa accgttgagc tcaggagggt ccgccgngag tacaaggcaa 60
 aaaagatctc tgaggaggaa tacgtcactg ccatcaagga ggaaatcaac aaggtcgtta 120
 agcttcaaga agagcttgac attgatgtgc ttgtgcatgg cgagcctgag agaaacgata 180
 tgggtgagta cttcggtgag cagctctccg gttttgcgtt cactgccaat ggctgggtgc 240
 agtattatgg gtcaaggngc gncaagctc cgatcatnga gggtgaccnc c 291

<210> 1075
 <211> 297
 <212> nucleic acid
 <213> Zea mays

 <400> 1075

 tgcacttgag nctttctggg atgggaagag caccgctgag gatctggaga aggttgctac 60
 cgacctcagg gccagtatct ggaagcagat ggctgangct gggatcaagt acatccccag 120
 caaacatttc tcgtactatg accaggctct tgacaccact gccatgctcg gcgctgtccc 180
 cgaacgttac tcatggactg gaggagagat tggatttgac acctacttct ccnnggccag 240
 gggcaacgcc actgttctctg ctatggagat gaccangtgg tttgacacca ataccac 297

<210> 1076
 <211> 312
 <212> nucleic acid
 <213> Zea mays

 <400> 1076

 acctatcatc gccgctgcgc cgtctcctcc gctcccgtcg tccggaaaga tggcgtccca 60
 cattgttgga taccctcgca tgggccccaa gagggagctc aagtttgac ttgagtcttt 120
 ctgggatggg aagagcaccg ctgaggatct ggagaagggt gctaccgacc tcagggccag 180
 tatctggaag cagatggctg atgctgggat caagtacatc ccagcaaca cttctcgta 240
 ctatgaccag gtccttgaca ccaactgccat gctcggcgct gtccccgaac gctactcatg 300

gactggagga ga

312

<210> 1077
<211> 307
<212> nucleic acid
<213> Zea mays

<400> 1077

gttgggaagg acaagottgt cgtatcaact tctgtctgc tcatgcacac cgctgtggat 60
cttgtgaacg agactaagct tgacagcgag attaatctt ggcttgcctt tgctgcccag 120
aagggtgttg aggggtganc tcttgctaag gcantggctg gtcaaaagga tgaggcttac 180
ttcgcagcaa atgtgtgtgc tcaggcctcg aggaaatcct caccocgtgt gaccaatgaa 240
gaagtccaga aggtgtgtgc tgctctcaag ggctctgacc accgccgtgg gaccaacgtt 300
agtgtcta 307

<210> 1078
<211> 274
<212> nucleic acid
<213> Zea mays

<400> 1078

ggcgtccac attgttggat accctcgcat gggccccaag anggagctca agtttgact 60
tgagtctttc tgggatggga agaacaccgc tgagnatctg gagaagggtg ctaccgacct 120
cagggccagt atctggaagc agatggctga tgctgggatc aagtacatcc ccagcaacac 180
tttctogtac tatgaccagg tcttgacac cactgccatg ctacgcgtg tccccgaacg 240
ttactcatgg actggaggag agattggatt tgac 274

<210> 1079
<211> 289
<212> nucleic acid
<213> Zea mays

<400> 1079

gagagattgg atttgacacc tacttctcca tggccagggg caacgccact gttcctgcta 60
tggagatgac caagtgggtt gacaccaact accactttat tgtccctgaa ttgggcccac 120
acaccaagtt ctctatgct tctcacaagg ctgttaacga atataaggag gctaaggcgc 180

tcggtgttga taccgtgcc a gtacttgttg gaccagtctc gtacctgttg ctctcaaagc 240
 ctgccaaagg tgtggagaag ggattccctc ttctttccct tcttagcag 289

<210> 1080
 <211> 286
 <212> nucleic acid
 <213> Zea mays

<400> 1080

ctaagottga cagcgagatt aagtcttggc ttgcttttgc tgcccagaag gttgttgagg 60
 ttgatgctct tgctaaggca ttggctggtc aaaaggatga ggcttacttc gcagcaaattg 120
 ctgcagctca ggccctcgagg aaatcctcac ccggtgtgac caatgaagaa gtccagaagg 180
 ctgctgctgc tctcaagggc tctgaccacc gccgtgagac caacgttagt gctagattgg 240
 atgccacgca gaagaagctt aacctcccca tcttccac cactac 286

<210> 1081
 <211> 323
 <212> nucleic acid
 <213> Zea mays

<400> 1081

cgccgctgag ccgtctctc cgctcccgtc gtccggaaag atggcgctcc acattgttgg 60
 atacctcgc atggggccca agagggagct caagtttgca cttgagtctt tctgggatgg 120
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctccaggcca gtatctggaa 180
 gcagatggct gatgctggga tcaagtacat cccagcaac accttctcgt actatgacca 240
 ggtccttgac accactgcc ngctcggcgc tgtcccgaa cgntactcat ggactgggng 300
 gagaaattgg gtttcacacc tat 323

<210> 1082
 <211> 305
 <212> nucleic acid
 <213> Zea mays

<400> 1082

cacaaaccgt tgcagctcag gaggggtccgc cgtgagtaca aggcaaaaaa gatctctgag 60
 gaggaatacg tcaactgcat caaggaggaa atcaacaagg tcgttaagct ccaagaagag 120

cttgacattg atgtgcttgt gcatggcgag cctgagagaa acgatatggg tgagtacttc 180
 ggtgagcagc tctccggttt tgcgttcact gccaatggct ggggtgcagtc ttatgggtca 240
 aggtgcgtca agcctccgat catctacggg gactgagccg cccaacccc atgaccgttt 300
 tctgg 305

<210> 1083
 <211> 281
 <212> nucleic acid
 <213> Zea mays
 <400> 1083

gtacaaggca aaaaatatct ctgaggagga atacgtcact gccatcaagg aggaaatcaa 60
 caaggctcgtt aagcnccaag aagagcttga cattgangtg cttgtgcatg gcgagcctga 120
 ganaaacgat atgggttgagt acttcgggtga gcagctctcc ggttttgctg tctactgcca 180
 tggttgggtg cagtcttatg ggtcaagggt cgtcaagcct ccgatcatct acggtgacgt 240
 gagccgcccc aaccccatga ccgttttctg gtcgnagacg g 281

<210> 1084
 <211> 314
 <212> nucleic acid
 <213> Zea mays
 <400> 1084

gaccagccga ggtttgagac ttgctaccag atcgctcttg cgatcaagaa ggaagtcgag 60
 gatctcgagg ctgggtggtat tcaggttatc caaatcgacg aggctgcact gagagaaggg 120
 ctgcgcctcc gcaaggctga gcatgcattc tacttggacn gggctgtcca cnccttcaga 180
 atcaccaacn gcgagatcca ggacaccacc cagatccaca cccacatgtg ctacnccaac 240
 ttcaacgaca tcatccactc gatcatcgac atggacgcgg ngtgatcacc acgaganncn 300
 ggtcgcgaga actg 314

<210> 1085
 <211> 281
 <212> nucleic acid
 <213> Zea mays
 <400> 1085

gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggtctt gaccaccgcc 60
 gtgggaccaa cgttagtgt agattggatg cccagcagaa gaagcttaac ctccccatcc 120
 ttcccaccac tacaatcggg tggttcccac aaaccgttga gctcaggagg gtccgccgtg 180
 agtacaagga aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca 240
 acaaggctgt taagcttcaa gaagagcttg acattgatgt g 281

<210> 1086
 <211> 306
 <212> nucleic acid
 <213> Zea mays

<400> 1086

ctatcatcgc cgctgcgcgc tctcctccgc tcccgctgtc cggaaagatg gcgtcccaca 60
 ttgttgata cctcgcgatg ggccccaaga gggagctcaa gtttgcaatt gagtctttct 120
 gggatgggaa gagcaccgct gaggatctgg agaagggtgc taccgacctc agggccagta 180
 tctggaagca gatggctgat gctgggatca agtacatccc cagcaacacc ttctcgtact 240
 atgaccaggt ccttgacacc actgccatgc tcggcgctgt ccccgaaacgc tactcatgga 300
 ctggag 306

<210> 1087
 <211> 265
 <212> nucleic acid
 <213> Zea mays

<400> 1087

tgctggtatt caggtgatcc agatcgatga ggcagctcta agggagggtc tgccactacg 60
 caagtcagag catgcattct acctggactg ggctgtccac tctttcagga tcaccaactg 120
 cggagtccag gacaccaccc agatccacac ccacatgtgc tactccaact tcaacgacat 180
 catccactcc atcatcgaca tggatgccga tgtgatcacg atcgagaact cccggtctga 240
 cgagaagcta ctgtccgtct tccgt 265

<210> 1088
 <211> 266
 <212> nucleic acid
 <213> Zea mays

<400> 1088

gcttaacctc cccatccttc ccaccactac aatcggttcg ttcccacaaa ccgttgagct 60

caggagggtc cgccgtgagt acaaggcaaa aaagatctct gaggaggaat acgtcactgc 120

catcaaggag gaaatcaaca aggtcgttaa gcttcaagaa gagcttgaca ttgatgtgct 180

tgtgcatggc gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg 240

ttttgcgttc actgccaatg gctggg 266

<210> 1089

<211> 311

<212> nucleic acid

<213> Zea mays

<400> 1089

agcagctctc cggttttgcg ttcactgcca atggctgggt gcagtcttat gggtaaggt 60

gcgtcaagcc tccgatcctc tacgggtgacg tgagccgcc caaccccatg accgttttct 120

ggtcgaagac ggacacagagc atgacgtctc gcccaatgaa ggggatgctg actggcccag 180

tcacaatcct caaotggctc ttcgtccgga atgaccagcc gaggtttgag acttgctacc 240

agatcgctct tgcgatcaag aaggaagtcg aggatttgag gctggtggta ttcaggttat 300

ccaaatcgac g 311

<210> 1090

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1090

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tgttgatac cctcgcatgg gcccgaagag ggagctcaag tttgacttg agtctttctg 120

ggatgggaag agcaccgctg aggatctgga gaaggttgct accgacctca gggccagtat 180

ctggaagcag atggctgatg ctgggatcaa gtacatcccc agcaacacct tctcgtacta 240

tgaccaggtc cttgacacca ctgccatgct cggcgtgtc cccgaacgct antcatggac 300

tggaggaga 309

<210> 1091

<211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 1091

 cgatcatoga catggacgcg gacgtgatca ccatcgagaa ctcgcggtcc gacgagaagc 60
 tgctctccgt gttccgtgag ggtgtcaagt acggcgcggg catcggtccc ggtgtctacg 120
 acatccactc ccccaggatc ccgtcggccg aggagatcgc cgaccgcatc gacaagatgc 180
 tggccgtgct ggacaccaac atcctctggg tgaacccga ctgcggcctc aagacccgca 240
 agtacacgga ggtcaagccc gccctgacca acatggtctc cgctgctaag t 291

<210> 1092
 <211> 295
 <212> nucleic acid
 <213> Zea mays

 <400> 1092

 tccgccgtga gtacaaggca aaaaagatct ctgaggagga atacgtcact gccatcaagg 60
 aggaaatcaa caaggctcgt ccagcttcaa gaancagctt gacattgatg tgcttgtgca 120
 tggcgagcct gagagaaaacg atatggttga gtacttcggt gagcagctct ccggttttgc 180
 gttcactgcc aatggctggg tgcagtctta tgggtcaagg tgcgtcaagc ctccgatcat 240
 ctacggtgac gtgagccgcc ccaaccccat gaccgttttc tggtcgaaga ggcac 295

<210> 1093
 <211> 285
 <212> nucleic acid
 <213> Zea mays

 <400> 1093

 ggtgcgtcaa gcctccgac atctacggtg acgtgagccg cccaacccc atgaccgttt 60
 tctggctgaa gaggcgcaga gcatgacgtc tcgcccaatg aaggggatgc tgactggccc 120
 agtcacaatc ctcaactggt ccttcgtccg gaatgaccag ccgaggtttg agacttgcta 180
 ccagatcgct cttgcgatca agaaggaagt cgaggatctc gaggctggtg gtattcaggt 240
 tatccaaatc gacgaggctg cactgagaga agggctgccg ctccg 285

<210> 1094

<211> 266
 <212> nucleic acid
 <213> Zea mays

 <400> 1094

 gctgcactga gagaagggct gccgctccgc aaggntgagc atgcattcta cttggactgg 60
 gntgtccact ccttcagaat tnccaaactgc gagatccagg acaccnccca gatccacacc 120
 cacatgtgct actccaactt caacgacatc atccactcga tcatcgacat ggacgaggac 180
 gtgatcacca ttgagaactc gcggtccgac ganaagctgc tctccgtgtt ccgcgagggc 240
 gtcaagtacg gcgcggggcat cggccc 266

<210> 1095
 <211> 327
 <212> nucleic acid
 <213> Zea mays

 <400> 1095

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 tgagacttgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc ttgaggctgg 120
 tggatttcag gttatccaaa tcgacgaggc tgcactgaga gaagggtgc cgctccgcaa 180
 ggctgagcat gcattctact tggactgggc tgtccactcc ttcagaatta ccaactgoga 240
 gatccaggac accacccaga tccacacca catgtgctat ccaattcacg acatcatcca 300
 togatcatcg ncatggngcg gcgtgat 327

<210> 1096
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 1096

 cagnacncac ccagatccac acccacatgt gctactccaa cttcaacgac atcatccact 60
 ccatcatcga catggatgcc gatgtgatca cgatcgagaa ctcccgtct gacgagaagc 120
 tactgtccgt cttccgtgag ggtgtgaagt acggagctgg cattggccct ggtgtctacg 180
 acatccactc tcttaggatt ccttcaccg aggagatcg agaccgcgtc gagaagatgc 240
 tcgngtgtg gananaant 259

<210> 1097
 <211> 280
 <212> nucleic acid
 <213> Zea mays

 <400> 1097

 cgacgagaag ctactgtccg tgttccgcga nggtgtgaag tacggagctg gaatcgggtcc 60
 aggtgtctac gacatccact cgcccaggat cccctccacc gaggagatcg cggaccgcgt 120
 caacaagatg ctgcgcgtgc tcgacaccaa catcctctgg gtgaaccctg actgcgggtct 180
 caagacaagc aagtacacgg aggtgaagcc tgccttgacc aacatggtct ccgccaccaa 240
 gtcctccgc acccagctcg gcagcgcgaa atgagtcaga 280

<210> 1098
 <211> 303
 <212> nucleic acid
 <213> Zea mays

 <400> 1098

 ggcgcctcct cacctatcat cgccgctgcg ccgtctcctc cgctcccgtc gtccggaaag 60
 atggcgctcc acattgttgg ataccctcgc atgggcccc aagaggagct caagtttgca 120
 cttgagtott tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac 180
 ctcagggcc a gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac 240
 actttctcgt actatgacca ggtccttgac accactgcc a tgctcggcgc tgtccccgaa 300
 cgt 303

<210> 1099
 <211> 305
 <212> nucleic acid
 <213> Zea mays

 <400> 1099

 cctcacctat catcgccgct gcgccgtctc ctccgctccc gtcgtccgga aagatggcgt 60
 cccacattgt tggataccct cgcattgggc ccaagaggga gctcaagttt gcacttgagt 120
 ctttctggga tgggaagagc accgctgagg atctggagaa ggttgctacc gacctcaggg 180
 ccagtatctg gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct 240

cgtactatga ccaggtcctt gacaccactg ccatgctcgg cgtgtcccc gaacgttact 300
catgg 305

<210> 1100
<211> 303
<212> nucleic acid
<213> Zea mays

<400> 1100

cngacgntgg ncgccgctgc gccgtctcct ccgtccccgt cgtccggaaa gatggcgctcc 60
cacattgttg gantaccctc gcatggggccc caagagggag ctcaagtttg cacttgagtc 120
ttttotgggat gggaagagca ccgtgagga tctggagaag gttgctaccg acctcagggc 180
cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca aactttctc 240
gtactatgac caggggcaac gccactgttc ctgctatgga gatgaccaag tggtttgaca 300
cca 303

<210> 1101
<211> 288
<212> nucleic acid
<213> Zea mays

<400> 1101

aggtgcgtca agcctccgat catctacggt gacgtgagcc gcccacccc catgaccgtt 60
ttctgggtcga agaggcacag agcatgacgt ctgcgccaat gaaggggatg ctgactggcc 120
cagtcacaat cctcaactgg tcttctgtcc ggaatgacca gccgaggttt gagacttgct 180
accagatcgc tcttgcgatc aagaaggaag tgcaggatct tgaggctggt ggtattcagg 240
ttatocaaat cgacgaggct gcactgagag aagggctgcc gctccgca 288

<210> 1102
<211> 275
<212> nucleic acid
<213> Zea mays

<400> 1102

cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgnc catggacgcg 60
gacgtgatca ccatcgagaa ctgcggtcc gacgagaagc tgctctccgt gttccgtgag 120

ggtgtcaagt acggcgcggg catcgggccc ggtgtctacg acatccactc cccagggatc 180
 ccgtcggccg aggagatcgc cgaccgcacg gacaagatgc tggccgtgct ggacaccaac 240
 atcctctggg tgaaccccgga ctgcggcctc aagac 275

<210> 1103
 <211> 319
 <212> nucleic acid
 <213> Zea mays

<400> 1103

gtgagccgcc ccaaccccat gaccgttttc tggtcgaaga cgggcgcaga gcatgacgtc 60
 tcgcccgaatg aaggggatgc tgactggccc agtcacaatc ctcaactggc cttcgtccg 120
 gaacgaccag ccgagggttg agacttgcta ccagatcgct cttgcgatca agaaggaagt 180
 cgaggatctt gaggttggtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 240
 agggctgccc ctccgcaagg ctgagcatgc attctacttg gatgggctgt ccatccttca 300
 gaataccaac tgcgagatc 319

<210> 1104
 <211> 277
 <212> nucleic acid
 <213> Zea mays

<400> 1104

ctcttgctaa ggcattggct ggtcaaaagg atgaggctta cttcgcagca aatgctgctg 60
 ctcaggcctc gaggaaatcc tcaccccgctg tgaccaatga agaagtccag aaggctgctg 120
 ctgctctcaa gggctctgac caccgcccgtg ggaccaacgt tagtgctaga ttggatgccc 180
 agcagaagaa gcttaacctc cccatccttc ccaccactac aatcggttcg tccccacaaa 240
 ccgttgagct caggaggggtc cgccgtgagt acaangc 277

<210> 1105
 <211> 308
 <212> nucleic acid
 <213> Zea mays

<400> 1105

caagaagagc ttgacattga tgtgcttgct catggcgagc ctgagagaaa cgatatgggtt 60

gagtacttcg gtgagcagct ctccggtttt gcgttcactg ccaatggctg ggtgcagtct 120
tatgggtcaa ggtgcgtcaa gctccgatac atctacgggtg angcnnaccg ccccaacccc 180
atgaccgttt tctggtcgaa gacggcacag agcatgacgt ctgcccgaat gaaggggatg 240
ctgactggcc cagtcacaat cctcaatggc ccttcgtccg gaatgaccan ccgangtttg 300
agatttct 308

<210> 1106
<211> 313
<212> nucleic acid
<213> Zea mays

<400> 1106

ggaggctaag gcgctcgggtg ttgataccgt gccagtactt gttggaccag tctcgtacct 60
gttgctctca aagcccgcca aggggtgtga gaagggattc cctcttcttt cccttcttag 120
cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggttctatg 180
gattcagttt gatgagccca ctcttgtcct cgaccttgat tctgacaaat tggttgcat 240
ctctgtgca taogcagaac ttgaatctgt actttctgga ttgaatgtgc ttgttgagac 300
ttactttgct gat 313

<210> 1107
<211> 279
<212> nucleic acid
<213> Zea mays

<400> 1107

gatgaggctt acttcgcagc aaatgctgca gctcaggcct cgaggaaatc ctcaccccgt 60
gtgaccaatg aagaagtcca gaaggctgct gctgctctca agggctctga ccaccgacct 120
gcgaccaacg ttagtgctag attggatgcc cagcagaaga agcttaacct cccatcctt 180
cccaccacta caatcgggtc gttcccacaa accgttgagc tcaggagggt ccgccgtgag 240
tacaaggcaa aaaagatctc tgaggaggaa tacgtcact 279

<210> 1108
<211> 264
<212> nucleic acid
<213> Zea mays

<400> 1108

aaacgaggag gtccaaaagg ctgcagctgc tttgaagggt tctgaccacc gccgtgctac 60

cactgtttgct gctagattgg atgctcagca gaaaaagctc aaccttcctg tccttccac 120

aaccacaatt ggttcattcc ctgagactgt ggaactcagg agggtcctgc gtgagtacaa 180

ggcaaagaag atctccgagg aggaatacat cagtgccatc aaggaagaaa tcagcaagggt 240

tgtcaagatc caagaggagc ttga 264

<210> 1109

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1109

ggttgatgct cttgctaagg cattggctgg tcaaaaggat gaggttact tcgcagcaaa 60

tgctgctgct caggcctcga ggaaatcctc acccctgtg accaatgaag aagtccagaa 120

ggctgctgct gctctcaagg gctctgacca ccgccgtggg accaacgtta gtgctagatt 180

ggatgccag cagaagaagc ttaacctccc catccttccc accactacaa tcggttcgtt 240

cccacaaacc gttgagctca ggagggtccg ccgtg 275

<210> 1110

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1110

ctcatggact ggaggagaga ttggatttga cacctacttc tccatggcca ggggcaacgc 60

cactgttctt gctatggaga tgaccaagtg gtttgacacc aactaccact ttattgtccc 120

tgaattgggc ccaaacacca agttctccta tgcttctcac aaggctgtta acgaatataa 180

ggaggctaag gcgctcgggtg ttgataccgt gccagtactt gttggaccag tctogtacct 240

gttgctctca aagcctgcc aaggtgtgga gaagggat 278

<210> 1111

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 1111
 cacctatcat cgccgctgcg ccgtctcttc cgctcccgtc gtccggaaaag atggcggtccc 60
 acattgtttg ataccctcgc atgggccccca agagggagct caagtttgca cttgagtctt 120
 tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca 180
 gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac accttctcgt 240
 actatgacca ggtccttgac accactgcca tgctcggcgc tgtccccana acgtactca 300
 tggactggag 310

<210> 1112
 <211> 265
 <212> nucleic acid
 <213> Zea mays

<400> 1112
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 gatgtgcttg tgcatggcga gcctgagaga aacgatatgg ttgagtactt cggtgagcag 120
 ctctccggtt ttgcgttcac tgccaatggc tgggtgcagt cttatgggtc aagggtgcgtc 180
 aagcctccga tcatctacgg tgacgtgagc cgcccccaacc ccatgaccgt tttctgggtc 240
 aagacggcgc agagcatgac gtctc 265

<210> 1113
 <211> 305
 <212> nucleic acid
 <213> Zea mays

<400> 1113
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 aggtgtctac gacatccact tcgcccagga tcccctccac cgaggagatc gcggaccgcg 120
 tcaacaagat gctcgcctg ctcgacacca acatcctctg ggtgaaccct gactgcggtc 180
 tcaagacacg caagtacacg gaggtgaagc ctgccctgac caacatggtc tccgccacca 240
 agctcatccg caccagctc ggcagcgcga aatgagtcag atttggtaga tccatgggtt 300
 gttga 305

<210> 1114

<211> 310
 <212> nucleic acid
 <213> Zea mays

 <400> 1114

 gtggtttgac accaactacc actttattgt cctgaattg ggcccaaaca ccaagttctc 60
 ctatgcttct cacaaggctg ttaacgaata taaggaggct aaggcgctcg gtgttgatac 120
 cgtgccagta cttgttggac cagtctcgta cctgttgctc tcaaagcctg ccaaggggtg 180
 ggagaaggga ttccctcttc ttcccttct tagcagnact tcccagtcta caaggaggtc 240
 attgctgagt tgaaggcagc tggggcttca ggattcagtt tgatgagccc atcttgtcct 300
 cgactngatt 310

<210> 1115
 <211> 331
 <212> nucleic acid
 <213> Zea mays

 <400> 1115

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 tatggttgag tacttcggtg agcagctctc cggttttgcg ttcaactgcca atggcnggg 120
 gcagtottat ggggtcaaggc gcgtaagcc tccgatcatc tacggtgacg tgagccgccc 180
 caaccccatg accgttttct ggtcgaagac gggcacagag catgacgtct cgcccaatga 240
 angggatgct gactggccca gtcacaatcc tcnactggtc ttcgtccgga atgaccagcc 300
 gangttgaga ttgctacnag atcgtctctg c 331

<210> 1116
 <211> 279
 <212> nucleic acid
 <213> Zea mays

 <400> 1116

 aacaaggctg ttaagcttca agaagagctt gacattgatg tgcttgtgca tggcgagcct 60
 gagagaaacg atatggttga gtacttcggt gagcagctct ccggttttgc gttcactgcc 120
 aatggctggg tgcagtctta tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac 180
 gtgagccgcc ccaaccccat gaccgttttc tggtcgaaga ggcacagagc atgacgtctc 240

gcccaatgaa ggggatgctg actggcccag tcacaatcc

279

<210> 1117
<211> 292
<212> nucleic acid
<213> Zea mays

<400> 1117

atcgccgctg cgccgtctcc tccgctcccg tcgtccggaa agatggcgtc ccacattggt 60
ggataccctc gcatggggcc ccaagaggga gctcaagttt gcacttgagt ctttctggga 120
tgggaagagc accgctgagg atctggagaa ggttgctacc gacctcaggg ccagtatctg 180
gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct cgtactatga 240
ccaggtcctt gacaccactg ccagtctcgg cgctgtcccc gaacgttact ca 292

<210> 1118
<211> 270
<212> nucleic acid
<213> Zea mays

<400> 1118

gaggaaatca acaaggctgt taagctccaa gaagagcttg acattgatgt gcttgtgcat 60
ggcgagcctg agagaaacga tatggttgag tacttcggtg agcagctctc cggttttgcg 120
ttcactgcca atggctgggt gcagtcttat gggtaaggt gcgtcaagcc tccgatcatc 180
tacggtgacg tgagccgccc caaccccatg accgttttct ggtcgaagag gcgcagagca 240
tgacgtctcg cccaatgaag gggatgctga 270

<210> 1119
<211> 252
<212> nucleic acid
<213> Zea mays

<400> 1119

cagcagaaga agcttaacct acccatcctt cccccacta caatagggtc gttcccacaa 60
accgttgagc tcaggagggt ccgccgtgag tacaaggcaa aaaagatctc tgaggaggaa 120
tacgtcactg ccatcaagga ggaaatcaac aaggctgcta agtccaaga agagcttgac 180
attgatgtgc ttgtgcatgg cgagcctgag agaaacgata tggttgagta cttcggtgag 240

cagctctccg gt

252

<210> 1120
<211> 284
<212> nucleic acid
<213> Zea mays

<400> 1120

cgttttctgg tcgaagaggg acagagcatg acgtctcgcc caatgaaggg gatgctgact 60
ggcccagtc caatcctcaa ctggtccttc gtccggaatg accagccgag gtttgagact 120
tgctaccaga togtctcttg gatcaagaag gaagtcgagg atcttgaggc tggtggtatt 180
caggttatcc aaatcgacga ggctgcactg agagaagggc tgccgctccg caaggctgag 240
catgcattct acttggaactg ggctgtccac tccttcagaa ttac 284

<210> 1121
<211> 287
<212> nucleic acid
<213> Zea mays

<400> 1121

cgctcatgca caccgctgtg gatcttgtga acgagactaa gcttgacagc gagattaagt 60
cttggettgc ttttgctgcc cagaagggtg ttgaggttga tgctcttgct aaggcattgg 120
ctgggtcaaaa ggatgaggct tacttcgcag caaatgctgc tgctcaggcc tcgaggaaat 180
cctcaccocg tgtgaccaat gaagaagtcc agaaggctgc tgctgctctc aagggctctg 240
accaccgcgc tgggaccaac gttagtgtga gattggatgc ccagcag 287

<210> 1122
<211> 290
<212> nucleic acid
<213> Zea mays

<400> 1122

cggacntggn ccgctgtgga tcttgtgaac gagactaagc ttgacagcga gattaagtct 60
tggtttgctt ttgctgccc gaaggttggt gaggttgatg ctcttgctaa ggcattggct 120
gggtcaaaagg atgaggctta ctctgcagca aatgctgcag ctcaggcctc gaggaaatcc 180
tcaccccggtg tgaccaatga agaagtccag aaggctgctg ctgctctcaa gggctctgac 240

caccgccgtg cgaccaacgt tagtgctaga ttggatgcc agcagaagaa 290

<210> 1123
 <211> 344
 <212> nucleic acid
 <213> Zea mays

<400> 1123

cgtaactgcc atcaaggagg naatcaacaa gtcgttaag ottcaagaag agcttgacat 60
 tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120
 gctctccggt ttgcggttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180
 caagcctccg atcancnacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240
 gaagacggca cagagcatga cgtctcgccc atgangggat gctgctggcc catcacaanc 300
 ctcaatngtc cttcgtccgg atgaccagcg agttgagntt ctac 344

<210> 1124
 <211> 261
 <212> nucleic acid
 <213> Zea mays

<400> 1124

cgtaactgcc atcaaggagg aaatcaacaa gtcgttaag ctcnaagaag agcttgacat 60
 tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120
 gctctccggt ttgcggttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180
 caagcctccg atcatctacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240
 gaagacggcg caaagcatga c 261

<210> 1125
 <211> 301
 <212> nucleic acid
 <213> Zea mays

<400> 1125

gcacgcgtgg gcgacatgga cgcggacgtg atcaccattg agaactcgcg gtccgacgag 60
 aagctgctct ccgtgttccg anngggcgtc aaacncggcg cgggcatcgg ccccggtgtc 120
 tacgacatcc actccccag gatcccgctg gctgaggaga tcgcccaccg catcgacaag 180

atgctggcgg tgctggacac caacatcctc tgggtgaacc ccgactgcgg cctcaagacc 240
cgcaagtaca cggaggtcaa gcccgccctg accaacaatgg tctccgccgc taagtcctc 300
c 301

<210> 1126
<211> 264
<212> nucleic acid
<213> Zea mays

<400> 1126

gggctgtcca ctcttcaga attaccaact gcnagatcca ggacaccacc cagatccaca 60
cccatatgtg ctactccaac ttcaacgnca tcatccactc gatcatcgac atggacgcgg 120
acgtgntcac cattgagaac tcgcgggtccg acgagaagct gctctccgtg ttccgcgagg 180
gcgtaagta cggcgcgggc atcggtcccg gtgtctacga catccnctcc ccagagatcc 240
cgtcggctga ggagatcgcc gacc 264

<210> 1127
<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1127

atcatccact cgatcatcga catggacgcg gacgtgatca ccattgagaa ctgcgggtcc 60
gacgagaagc tgctctccgt gttccgcgag ggcgtcaagt acggcgcggg catcggtccc 120
ggtgtctacg acatccactc cccagagatc ccgtcggctg aggagatcgc cgaccgcctc 180
gacaagatgc tggcgggtgct ggacaccaac atcctctggg tgaaccccga ctgcggcctc 240
aagacccgca agtacacgga ggtcaagccc gcctgac 277

<210> 1128
<211> 246
<212> nucleic acid
<213> Zea mays

<400> 1128

gggctgtcca ctctttcagg atcaccaact gcggagtcca ggacaccacc cagatccaca 60
cccatatgtg ctactccaac ttcaacggac atcatccact ccatcatcga catggatgcc 120

gatgtgatca cgatcgagaa ctcccgatct gacgagaagc tactgtccgt cttccgtgag 180
 ggtgtgaagt acggagctgg cattggccct ggtgtctacg acatccactc tcctaggatt 240
 ccctcc 246

<210> 1129
 <211> 270
 <212> nucleic acid
 <213> Zea mays
 <400> 1129

aaatcaacaa ggtcgttaag ctccaagaag agcttgacat tgatgtgctt gtgcatggcg 60
 agcctgagag aaacgctatg gttgagtact tcggtgagca gctctccggt tttgcgttca 120
 ctgccaatgg ctgggtgcag tcttatgggt caaggtgctg caagcctccg atcatctacg 180
 gtgacgtgag ccgccccaac cccatgaccg tttcttggtc gaagaggcgc agagcatgac 240
 gtctcgccca atgaagggga tgctgactgg 270

<210> 1130
 <211> 492
 <212> nucleic acid
 <213> Zea mays
 <400> 1130

ttannttccg cgccagtaag gtaagaattc ccgggncgac ccacgcgtca gtgagaactc 60
 gcggtccgac gagaagctgc tctccgtggt ccgcgagggc gtcaagtacg gcgcgggcat 120
 cgccccgggt gtctacgaca tccactcccc caggatcccc tcggctgagg agatcgccga 180
 ccgcatcgac aagatgctgg cgggtgctgga caccaacatc ctctgggtga accccgactg 240
 cggcctcaag acccgcaagt acacggaggt caagcccgcc ctgaccaaca tgggtctccgc 300
 cgctaaagct catccgcacc cagctcgcca gcgccaagtg agcacctttt ttttgccctt 360
 ttcgtttccg aaganggcgt cntcgatgcc aatttgtttc caataaacan ggctgtnncc 420
 cgccgttccg gttgtnancc gttcgggtgg aaggttanna attttcctga atttcccccc 480
 aanggcggtt ng 492

<210> 1131
 <211> 250
 <212> nucleic acid

<213> Zea mays

<400> 1131

aactcaggag ggttcgccgt gaatacaagg cnaagaagat caccgaggac gaatacatca 60
 gtgccatcaa ggaagaaatc agcaagggtt tcaagatcca agaggagctt gacattgatg 120
 tgcttgtgca tggagagcca gagagaaatg acatgggttg gtacttcggt gncnaattat 180
 ctggttttgc gttcaactgcc aacggatggg tgcaatccta tggatcacgc tgtgtgaagc 240
 cacccattat 250

<210> 1132

<211> 262

<212> nucleic acid

<213> Zea mays

<400> 1132

tottggcttg cttttgctgc ccagaagggt gttgaggttg atgctcttgc taaggcattg 60
 gctgggtcaa aggatgagge ttacttcgca gcaaagtctg ctgctcaggc ctcgaggaaa 120
 tcctcacccc gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggtctt 180
 gaccaccgcc gtgggaccaa cgttagtgtc agattggatg cccagcagaa gaagcttaac 240
 ctccccatcc ttcccaccac ta 262

<210> 1133

<211> 269

<212> nucleic acid

<213> Zea mays

<400> 1133

gctgctcagg cctcgaggaa atcctcaccc cgtgtgacca atgaagaagt ccagaaggct 60
 gctgctgctc tcaagggtc tgaccaccgc cgtgcgacca acgttagtgc tagattggat 120
 gccagcaga agaagcttaa cctccccatc cttcccacca ctacaatcg ttcgttcna 180
 caaacggttg agctcaggag ggtccgccgt gagtacaagg caaaaaagat ctctgaggag 240
 gaatacgtca ctgccatcaa ggaggaaat 269

<210> 1134

<211> 269

<212> nucleic acid

<213> Zea mays

<400> 1134

cgccgctgcg ccgtctcctc cgctcccgtc gtcgggaaag atggcgctcc acattgttgg 60
 ataccctcgc atggggcccca agagggagct caagtttgca cttgagtctt tctgggatgg 120
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180
 gcagatggct gatgctggga tcaagtacat cccagcaac acccttctgt actatgacca 240
 ggtccttgac accactgcc a tgctcggcg 269

<210> 1135

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1135

gcgatcaaga aggaagtcga ggatcttgag gctggtggta ttcaggttat ccaaatcgac 60
 gaggctgcac tgagagaagg gctgccgntc cgcaaggctg agcatgcatt ctacttggac 120
 tgggctgtcc actccttcag aattaccaac tgcgagatcc aggacaccac ccagatccac 180
 acccacatgt gctantccaa cttcaacgac atcatccact cgatcatcga catggacgng 240
 gacgtgatna ccattgagaa ntcgggtccg angggnagct gctctcngtg ttcngggag 299

<210> 1136

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1136

ctcaccctgt gtgaccaatg aagaagtcca gaaggctgct gctgctctca agggctctga 60
 ccaccgctgt gggaccaacg ttagtgctag attggatgcc cagcagaaga agcttaacct 120
 ccccatcctt cccaccacta caatcggttc gttcccacaa accgttgagc tcaggagggt 180
 ccgccgtgag tacaaggcaa aaaagatctc tgaggaggaa tacgtcactg ccatcaagga 240
 ggaaatcaac aaggtcgtta agcttcaaga agagct 276

<210> 1137

<211> 257

<212> nucleic acid

<213> Zea mays

<400> 1137

ccaactgcga gatccaggac accaccaga tccacacca catgtgctac tccaacttca 60
acgacatcat ccaactgata atcgacatgg acgcggacgt gatcaccatc gagaactcgc 120
ggtcgcacga gaagctgctc tccgtgttcc gtgagggtgt caagtacggc gcggggcatcg 180
gccccgggtgt ctacgacatc cactccccc ggatcccgtc ggccgaggag atcgccgacc 240
gcatcgacaa gatgctg 257

<210> 1138

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1138

gcttacttgc cagcaaatgc tgctgctcag gcctcgagga aatcctcacc ccgtgtgacc 60
aatgaagaag tccagaaggc tgctgctgct ctcaagggct ctgaccaccg ccgtgggacc 120
aacgttagtg ctagattgga tgcccagcag aagaagctta acctcccat ccttcccacc 180
actacaatgc gttcgttccc acaaaccgtt gagctcagga gggccgccc tgagtacaag 240
gcaaaaaaga tctctgagga ggaatacgtc actgc 275

<210> 1139

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 1139

agatggcaca gagcatgacc cctcgtccca tgaagggaat gttgactggc ccggtcacao 60
tctgaactg gtcattcgtc aggaacgacc agcctaggtt tgagacatgc taccaaatag 120
ctcttgcaat caaaaaggag gttgaggatc ttgaggctgc tggattcag gtgatccaga 180
togatgaggc agctctaagg gagggctctgc cactacgcaa gtcagagcat gcattctacc 240
tggactgggc tgtccactct ttcaggatca ccaactgcgg agtccaggac accac 295

<210> 1140

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 1140

gcattctctg ctgcatacgc agaacttgaa tctgtacttt ctggattgaa tgtgcttggt 60
gagacttact ttgctgatgt tctgtctgag tctacaaga ccctaacatc cctgagcagt 120
gtgactgctt atgggttttga tcttgctcgt ggaacccaaa ctcttgggct tgtcacgagt 180
gctgggttcc ctgctggaaa gtacctcttt gctgggtgtg tggatggacg caacatctgg 240
gctgatgata ttgctacatc tctcagcact ctccagtctc ttgaggctgt tgttgggaag 300
gacaagcttg tcgtat 316

<210> 1141

<211> 445

<212> nucleic acid

<213> Zea mays

<400> 1141

ccactcttgt cctcgacctt gattctgaca aattggctgc attctctgct gcatacgcag 60
aacttgaatc tgtactttct ggattgaatg tgcttggtga gacttacttt gctgatgttc 120
ctgctgagtc ctacaagacc ctaacatctc tgagcagtgt gactgcctat ggttttgata 180
ttgtccgtgg aacccaaact cttgggcttg tcacgagtgc tggtttccct gctggaaagt 240
acctctttgc tgggtgtgtg gatggacgca acatctgggc tgatgatctt gctacatctc 300
taacactctt cagtctcttg aggctgttgt tgggaangac aagcttgctg atcaacttnc 360
tgctcgctca tgcacaccgn tgtggatctt gngaaccaga ctnagcttga cagcagatna 420
atctggcttg ctttgctgcc anaag 445

<210> 1142

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 1142

gccaaagacg ttgttgaggt tgatgctctt gctaaggcat tggctggtca aaaggatgag 60
gcttacttcg cagcaaattgc tgcagctcag gcctcgagga aatcctcacc ccgtgtgacc 120
aatgaagaag tccagaaggc tgctgctgct ctcaagggt ctgaccaccg ccgtgcgacc 180

aacgtagtg ctagattgga tgcccagcag aagaagctta acctcccat cttcccacca 240
tacaatcggg tcgttccac aaaccgttga gtcaggagg gtccgccgtg agtacaaggc 300
a 301

<210> 1143
<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1143

gtggtattca gggtatccaa atcgacgagg ctgcactgag agaagggctg ccgctccgca 60
aggctgagca tgcattctac ttggactggg ctgtccaact cttcagaatc accaactgcg 120
agatccagga caccaccng atccacaccc acatgtgcta ctccaacttc aacgacatca 180
tccactgat catcgacatg gacgaggacg tgatcaccat cgagaatcgc ggtccgacga 240
gaagctgtc tccgtgttcc gtgaggtgtc aagtacg 277

<210> 1144
<211> 281
<212> nucleic acid
<213> Zea mays

<400> 1144

cgccgtgcg ccgtctctc cgtcccgtc gtccggaaag atggcgtccc acattgttgg 60
ataccctcgc atggggccca agagggagct caagtttgca cttgagtctt tctgggatgg 120
gaagagcacc gctgaggatc tggagaangt tgctaccgac ctccaggcca gtatctggaa 180
gcagatggct gatgctggga tcaagtacat cccagcaac actttctcgt actatgacca 240
ggtcctttga caccactgcc atgctcggcg ctgtccccga a 281

<210> 1145
<211> 247
<212> nucleic acid
<213> Zea mays

<400> 1145

gccgtgctac cactgttgct gctagattgg atgctcagca gaaaaagctc aaccttctg 60
tccttccac aaccacaatt ggttcattcc ctgagactgt ggaactcagg aggtccgctc 120

gtgagtacaa ggcaaagaag atctccgagg aggaatacat cagtgccatc aaggaagaaa 180
 tcagcaaggt tgtcaagatc caagaggagc ttgacattga tgtgcttgtg catggagagc 240
 cagagag 247

<210> 1146
 <211> 268
 <212> nucleic acid
 <213> Zea mays
 <400> 1146

tgctgactgg ccagtcaca atcctcaact ggtccttcgt ccggaatgac cagccgaggt 60
 ttgagacttg ctaccagatc gctcttgcca tcaagaagga agtcgaggat cttgaggctg 120
 gtggtattca ggttatccaa atcgacgagg ctgcaactgag agaagggctg ccgctccgca 180
 aggctgagca tgcattotac ttggactggg ctgtccactc cttcagaatt accaactgcg 240
 agatccagga caccacccag atccacac 268

<210> 1147
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 1147

cggnacggtg gcnacctatc atcgccgctg cgccgtctcc tccgctcccg tcgtccggaa 60
 agatggcgtc ccacattggt ggataccctc gnntgggccc caagagggag ctcaagtttg 120
 cacttgagtc tttctgggat ggggaagagca ccgctgagga tctggagaag gttgctaccg 180
 acctcagggc cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 240
 acactttctc gtactatgac caggtccttg acaccactgc catgctcggc 290

<210> 1148
 <211> 269
 <212> nucleic acid
 <213> Zea mays
 <400> 1148

caaggagatc atcgcccagg gtgacaaacg aggaggtcca gaaggctgca gctgctttga 60
 ggggatctga ccaccgccgt tctaccactg tttctgctag attggatgct cagcagaaaa 120

agctcaacct tctgtcctt cccacaacca caattggttc attccctcag actgtggaac 180
 tcaggagggt tcgccgtgaa tacaaggcaa agaagatcac cgaggacgaa tacatcagtg 240
 ccatcaagga agaaatcagc aaggttgtc 269

<210> 1149
 <211> 280
 <212> nucleic acid
 <213> Zea mays

<400> 1149

caagaagagc ttgacatnna tgtnccttgtn catngcgagc ctgagagaaa cgatatggtt 60
 gagnacntcg gtgagcagct cncgggtttt gcgttcacng ccaatggctg ggtgcagtct 120
 tatgggtcaa ggtgogtcaa gcctccgac atctacgggtg acgtgagcng ccccaacccc 180
 atgaccgttt tnggtcgaa gacngcgag agcatgacgt ctcgcccaat gaaggggatg 240
 ctganggccc agtcacaatc ctcaacnggt ccttcgtccg 280

<210> 1150
 <211> 323
 <212> nucleic acid
 <213> Zea mays

<400> 1150

gagaactcgc ggtccgacga gaagctgctc tccgtgttcc gtgaggggtg caagtacggc 60
 gcgggcatcg gccccggtgt ctacgacatc cactcccca ggatcccgtc ggccgaggag 120
 atcgccgacc gcacgacaa gatgctggcc gtgctggaca ccaacatcct ctgggtgaac 180
 cccgactgcg gcctcaagac ccgcaagtac acggaggtea agcccgccct gaccaacatg 240
 gtctccgctg ctaagtcatt cgcaccacgc tcgccagcgc caagtgagca ctttttttgc 300
 ctttttctgt tccgaggagg ggt 323

<210> 1151
 <211> 325
 <212> nucleic acid
 <213> Zea mays

<400> 1151

caccacccac atctcagttc accccggcgt cgtcctcttc ccccggcgt actctccgc 60

tccacgttcc aaaggaaaga tggcgtccca tattgttga taccctcgca tgggccccaa 120
gagagagctc aagtttgcct tggagtcttt ctgggatggg aagagcagcg ccgatgattt 180
agagaaagtt gcctctgacc tcaggtctag catctggaag caaatgtcag aagctgggat 240
caagtacatt cccagcaaca ccttctcgta ctatgaccag gtccttgata ccacggccat 300
gottggtgtg tcccagagcg ctact 325

<210> 1152
<211> 272
<212> nucleic acid
<213> Zea mays

<400> 1152

ggacgtggac ttgcgatggg cccaagagg gagctcaagt ttgcacttga gtcantctgg 60
gatgggaaga gcaccgctga ggatctggag aaggttgcta ccgacctcag ggccagtatc 120
tggangcaga tggctgatgc tgggatcaag tacatcccca gcaacacctt ctctactat 180
gaccaggtcc ttgacaccac tgccatgctc ggcgctgtcc ccgaacgcta ctcatggact 240
ggaggagaga ttgggtttga cacctacttc tc 272

<210> 1153
<211> 232
<212> nucleic acid
<213> Zea mays

<400> 1153

gggctgtcca ctcttcaga atcaccaact gcgagatcca ggacaccacc cagatccaca 60
cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac atggacgcgg 120
acgtgatcac catcgagaac tcgcggtccg acgagaagct gctctccgtg ttccgtgagg 180
gtgtcaagta cggcgcgggc atcggtcccg gtgtctacga catccactcc cc 232

<210> 1154
<211> 285
<212> nucleic acid
<213> Zea mays

<400> 1154

atcgacgagg ctgcactgag agaagggctg ccgctccgca aggctgagca tgcattctac 60

ttggactggg ctgtccactc cttcagaatc accaactgcg aaatccagga caccacccag 120
atccacaccc acatgtgcta ctccacttca acgacatcat ccaactcgatt nccgacatgg 180
acgcggacgt gatcaccatc gagaactcgc ggtccgacga gaagctgctc tccgtgttcc 240
gtgaggtgtc aagtacggcg cgggcacgcg ccccggtgtc tacga 285

<210> 1155
<211> 327
<212> nucleic acid
<213> Zea mays

<400> 1155

cccggtctga cgagaagcta ctgtccgtct tccgtgaggg tgtgaagtac ggagctggca 60
ttggccctgg tgtctacgac atccacttct cctaggattc cctccaccga ggagatcgca 120
gaccgcgtcg agaagatgct cgcggtgctc gacaccaaca tcctctgggt gaaccctgac 180
tgtggtctca agacacgcaa gtacacggag gtcaagcccg ccctgaccaa catggtctcg 240
gccaccaagc tcattccgac ccagcttgcc agcgcgaaat gagtcgtttg atagtccatg 300
gtctgatagc gcggaatgag ccagttt 327

<210> 1156
<211> 284
<212> nucleic acid
<213> Zea mays

<400> 1156

cgatgacacc attgaganct cgcggtccga cgagaagctg ctctccgtgt tccgcgaggn 60
gcgtcaagta cggcgcgggc atcgccccg gtgtctacna catccactcc ccaggatcc 120
cgtcggctga ggagatcgcc gaccgcatcg acaagatgct ggcggtgctg gacaccaaca 180
tcctctgggt gaaccccgac tgcggcctca agacccgcaa gtacacggag gtcaagcccg 240
ccctgaccaa catggtctcc gccgctaagc tcattccgac ccag 284

<210> 1157
<211> 293
<212> nucleic acid
<213> Zea mays

<400> 1157

gtcttgggct tgcttttgct gcccagaagg ttgttgaggt tgatgctctt gctaaggcat 60
 tgggctgggc aaaaggatga ggcttacttc gcagcaaagt ctgctgctca ggctctgagg 120
 aaatcctcac cccgtgtgac caatgaagaa gtccagaagg ctgctgctgc tctcaagggc 180
 totgaccacc gccgtgggac caacgttagt gctagattgg atgccagca gaagaagctt 240
 aacctcccca tcttccac cactacaatc ggttcgttcc cacaaaccgt tga 293

<210> 1158
 <211> 309
 <212> nucleic acid
 <213> Zea mays

<400> 1158

cgccgctgcg ccgtctctc cgctcccgtc gtccggaag atggcgnccc acattgttgg 60
 ataccctcgc atgggccccca agaggagct caagtttgca cttgagtctt tctgggatgg 120
 gaagggcacc gctgaggatc tggagaaggt tgctaccgac ctccaggcca gtatctggaa 180
 gcagatggct gatgctggga tcaagtacat cccagaanc actttctcgt actatgacca 240
 agtcttgaca ccattgccatg ctccggcgtg tccccgaac ttactcatgg actggaggag 300
 agatggatt 309

<210> 1159
 <211> 462
 <212> nucleic acid
 <213> Zea mays

<400> 1159

cgtgcggtgn tccgcgaggg cgtcaagtac ggcgcnggca tcggccccgg tgtctacgac 60
 ntccactccc ccaggatccc gtcggctgag gagatcgccg accgcatcga caagatgctg 120
 gcggtgctgg acaccaacat cctctgggtg aaccccgact gcggcctcaa gaccgcaag 180
 tacacggagg tcaagccgc cctgaccaac atgggtctccg ccgctaagct catccgcacc 240
 cagctcgcca gcgccaagtg agcacctttt ttttgctttt ttcgtttccg aggagggccg 300
 tcgtcgatgc caatttgttt ccaataaaca gggcttgctg ccgccgttct gttgtactcc 360
 gtctgtggtt aggttaagta gttttcttga tctcgcccc acgcggtacc tgnnttacta 420
 ctctctggtt ggtgggttct gaagcaaagt tnnccgttta ct 462

<210> 1160
 <211> 231
 <212> nucleic acid
 <213> Zea mays

 <400> 1160

 ggctgcactg agagaagggc tgccgctccg caaggctgag catgcattct acttggactg 60
 ggctgtccac tccttcagaa tcaccaactg cgagatccag gacaccaccc agatccacac 120
 ccacatgtgc tactccaact tcaacgacat catccactcg atcatcgaca tggacgcgga 180
 cgtgatcacc atcgagaact cgcgggtccga cgagaagctg ctctccgtgt t 231

<210> 1161
 <211> 275
 <212> nucleic acid
 <213> Zea mays

 <400> 1161

 gttgactggt ccggtcacaa tcctganctg gtcattcgtc aggaacgacc agcctagggt 60
 tgagacatgc taccaaatag ctcttgcaat caaaaaggag gttgaggatc ttgaggcnnc 120
 nggtattcag gtgatccaga tcgatgaggc agctctaagg gagggctctgc cactacgcaa 180
 gtcagagcat gcattctacc tggactgggc tgtccactct ttcaggatca ccaactgcgg 240
 agtccaggac accaccaga tccacacca catgt 275

<210> 1162
 <211> 290
 <212> nucleic acid
 <213> Zea mays

 <400> 1162

 gagagattgg atttgacacc tacttctcca tggccagggg caacgccact gttcctgcta 60
 tggagatgac caagtggttt gacaccaact accactttan tgtccctgga nttgggcca 120
 aacaccaagt tctcctatgc ttctcacaag gctgttaacg aatataagga ggctaaggcg 180
 ctcggtgttg ataccgtgcc agtacttggt ggaccagtct cgtacctgtt gctctcaaag 240
 cctgccaaagg gtgtggagaa gggattcnct cttctttccc ttcttagcag 290

<210> 1163

<211> 342
 <212> nucleic acid
 <213> Zea mays

 <400> 1163

 cngtgacgtg agccgcccc accccatgac cgttttctgg tcgaagacgg cgcagagcat 60
 gacgtctcng ccaatgaaag gngatgctga ctggcccagt ncacaatcct caactgggtcc 120
 ttcgtcogga atgaccagcc gaggtttgag acttgctacc agatcgctct tgcgatcaag 180
 aaggaagtcg aggatctcga ggctgggtgg attcagggtta tccaaatcga cgaggctgca 240
 ctgagagaag ggctgccgct ccgcaaggct gagcatgcat ctacttgga tgggctgtcc 300
 atccttcaga atcaccaatg cgnagaccag gacaccaccc ag 342

<210> 1164
 <211> 264
 <212> nucleic acid
 <213> Zea mays

 <400> 1164

 aaccccatga ccgttttctg gtcgaagagg cacagagcat gacgtctcgc ccaatgaagg 60
 ggatgctgac tggcccagtc acaatcctca actggtcctt cgtccggaat gaccagccga 120
 ggtttgagac ttgctaccag atcgctcttg cgatcaagaa ggaagtcgag gatcttgagg 180
 ctggtggtat tcaggttatc caaatcgacg aggctgcact gagagaaggg ctgccgctcc 240
 gcaaggctga gcatgcattc tact 264

<210> 1165
 <211> 298
 <212> nucleic acid
 <213> Zea mays

 <400> 1165

 gtccagaagg ctgcagctgc tttgagggga tctgaccacc gccgttctac cactgtttct 60
 gctagattgg atgctcagca gacaaagctc aaccttnctg tccttccac aaccacaatt 120
 gnttcattcc ctcagactgt ggaactcagg agggttcgcc gtgaatacaa ggcaaagaag 180
 atcaccgagg acgaaaanat cagtgccatc aaggaagaaa ttcagcaagg ttgtcaagat 240
 ccaagaggag ntgacattga tgtgcttgtg catggagagc cagagagaat gacatggt 298

<210> 1166
 <211> 226
 <212> nucleic acid
 <213> Zea mays

 <400> 1166

 gcttgacatt gatgtgcttg tgcattggaga gccagagaga aatgacatgg ttgagtactt 60
 cgggtgagcaa ttatctgggtt ttgcgttcac tgccaacgga tgggtgcaat cctatggatc 120
 acgctgtgtg aagccaccca ttatctacgg tgatgtcagc cggccgaacc ccatgactgt 180
 tttctgggtcc aagatggcac agagcatgac cctcgtccc atgaag 226

<210> 1167
 <211> 254
 <212> nucleic acid
 <213> Zea mays

 <400> 1167

 gtccggaatg accagccgag gtttgagact tgctaccaga tcgctcttgc gatcaagaag 60
 gaagtogagg atcttgaggc tgggtgtatt caggttatcc aaatcgacga ggctgcactg 120
 agagaagggc tgccgctccg caaggctgag catgcattct acttggactg ggctgtccac 180
 tccttcagaa ttaccaactg cgagatccag gacaccaccc agatccacac ccacatgtgc 240
 tactccaact tcaa 254

<210> 1168
 <211> 251
 <212> nucleic acid
 <213> Zea mays

 <400> 1168

 ctccggtttt gcgttcaactg ccaatggctg ggtgcagtct tatgggtcaa ggtgcgtcaa 60
 gcctccgac atctacggtg acgtgagccg cccaacccc atgaccgttt tctggtcgaa 120
 gaggcgcaga gcatgacgtc tcgcccgaatg aaggggatgc tgactggccc agtcacaatc 180
 ctcaactgggt ccttcgtccg gaatgaccag ccgaggtttg agacttgcta ccagatcgct 240
 cttgcgatca a 251

<210> 1169

<211> 279
 <212> nucleic acid
 <213> Zea mays

 <400> 1169

 atcgccgctg cgccgtctcc tccgctcccg tcgtncggaa agatggcgtc ccacattggt 60
 ggataccctc gcatggggccc caagagggag ctcaagtttg cacttgagtc tttctgggat 120
 gggaagagca ccgctgagga tctggagaag gttgcnacng aacttnaggg ccagtatctg 180
 gaagcagang gctgatgctg ggatcaagta catccccagc aacactttct cgtactatga 240
 ccaggtcctt gacacnactg ccatgctggg gctggtccc 279

<210> 1170
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 1170

 atgggtcaag gtgcgtaag cctccgatca tctacggtga cgtgagcngc cccaacccca 60
 tgaccgtttt ctgggtgaag agggcgagan catgacgtct cgcccaatga aggggatgct 120
 gactggccca gtcacaatcc tcaactggtc cttcgctccg aatgaccagc cgaggtttga 180
 gaattgctac cagatcgctc ttgcgatcaa gaaggaagtc gaggatctcg aggctggtgg 240
 tattcagggt atccaaatcg acgaggct 268

<210> 1171
 <211> 90
 <212> nucleic acid
 <213> Zea mays

 <400> 1171

 cgtcggctga ggagatcgcc gaccgcactn acaagatgct ggcggtgctg gacaccaaca 60
 tcntctgggt gaaccccgac tgcngcctca 90

<210> 1172
 <211> 254
 <212> nucleic acid
 <213> Zea mays

 <400> 1172

gtcctccgct cccgtcgtcc ggaaagatgg cgtcccacat tgttggatac cctcgcatgg 60
 gcccgaagag ggagctcaag tttgcacttg agtctttctg ggatgggaag agcaccgctg 120
 aggatctgga gaaggttget accgacctca gggccagtat ctggaagcag atggctgatg 180
 ctgggatcaa gtacatcccc agcaacacct tntcgtacta tgaccaggtc cttgacacca 240
 tgccatgctc ggnt 254

<210> 1173
 <211> 251
 <212> nucleic acid
 <213> Zea mays

<400> 1173

caaacgagga ggtccagaag gctgcagctg ctttgagggg atctgaccac cgccgttcta 60
 ccaactgtttc tgctagattg gatgctcagc agaaaaagct caaccttcct gtccttccca 120
 caaccacaat tggttcattc cctcagactg tggaactcag gagggttcgc cgtgaatata 180
 aggcaaagaa gatcaccgag gacgaatata tcagtgncat caaggaagaa atcagcaagg 240
 ttgtcaagat c 251

<210> 1174
 <211> 246
 <212> nucleic acid
 <213> Zea mays

<400> 1174

ccgatgtgat cacgatcgag aactcccggt ctgacgagaa gctactgtcc gtcttccgtg 60
 aggggtgtgaa gtacggagct ggcattggcc ctgggtgtcta cgacatccac tctcctagga 120
 ttccctccac cgaggagatc gcagaccgcg tcgagaagat gctcgccgtg ctcgacacca 180
 acatcctctg ggtgaacctt gactgtggtc tcaagacacg caagtacacg gaggtcaagc 240
 ccgccc 246

<210> 1175
 <211> 258
 <212> nucleic acid
 <213> Zea mays

<400> 1175

gaacgatcag cctaggtttg agacatgcta ccaaatagct cttgcaataa aaaaggaggt 60
 tgaggatctt gaggtgctg gtattcaggt gatccagatt gatgaggcag ctctaaggga 120
 ggggtetgcca ctgcgcaagt cagagcatgc attctacctg gactgggctg tccactcttt 180
 cagaatcacc aactgcggag tccaggacac caccagatc cacaccata tgtgctactc 240
 caacttcaac gacatcat 258

<210> 1176
 <211> 276
 <212> nucleic acid
 <213> Zea mays

<400> 1176

cggacontggn nagnngnaaat cctcaccocg tgtgaccaat gaagaagtcc agaaggctgc 60
 tgctgtcttc aagggtcttn aanaccgocg nngnaccaac gttagtgtta gattggatgc 120
 ccagcagaag aagcttaacc tccccatcct tcccaccact acaatcggtt cgttcccaca 180
 aaccgttgag ctgaggaggg tccgccgtga gtacaaggca aaaaagatct ctgaggagga 240
 atacgtcact gccatcaagg aggaaatcaa caaggt 276

<210> 1177
 <211> 297
 <212> nucleic acid
 <213> Zea mays

<400> 1177

acgagtgtctg gtttccctgc tggaaagtac ctctttgctg gtgttggtga tggacgcaac 60
 atctgggctg atgatcttgc tacatctctc agcactctcc agtctcttga ggctgttggt 120
 gggaaggaca agcttgctgt atcaacttcc tgctcgctca tgcacaccgc tgtggatctt 180
 gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgctttttg cccagaaggt 240
 tgttgagggt gatgtcttgc ctaaggcatt ggctgggcca aaagatgagc ttacttc 297

<210> 1178
 <211> 283
 <212> nucleic acid
 <213> Zea mays

<400> 1178

ggaggctaag gcgctcgggtg ttgataccgt gccagtactt gttggaccag tctcgtacct 60
gttgcctctca aagcctgccca aggggtgtgga gaagggattc cctcttcttt cccttcttag 120
cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggcttcatg 180
gattcagttt gatgagccca ctcttgtcct cgaccttgat tctgacaaat tggtctgcatt 240
ctctgctgca tacgcagaac ttgaatctgt actttctgga ttg 283

<210> 1179
<211> 84
<212> nucleic acid
<213> Zea mays

<400> 1179

ancttctngt actatgacca ggtccttgan accagtncca tgctcggcgc tgtccccgaa 60
cgctaccatg gactggacga gaga 84

<210> 1180
<211> 315
<212> nucleic acid
<213> Zea mays

<400> 1180

gtttgatgag cccactcttg tctctgacct tgattctgac aaattggctg cattctctgc 60
tgcatacgca gaacttgaat ctgtactttc tggattgaat gtgcttggtg agacttactt 120
tgctgatgtt cctgctgagt cctacaagnc cctaacatcc ctgagcagtg tgactgctta 180
tggttttgat cttgtccgtg gaacccaaac tcttgggctt gtcacgagtg ctggtttccc 240
tgctggaaaag tacctctttg ctggtgttgt ggatggacgc aacatctggg ctgatgatct 300
tgctacatct ctgag 315

<210> 1181
<211> 267
<212> nucleic acid
<213> Zea mays

<400> 1181

agccgccccca acccatgac cgttttctgg tcgaagaggc gcagagcatg acgtctcgcc 60
caatgaaggg gatgctgact ggcccagtca caatcctcaa ctggctcctt gtccggaatg 120

accagccgag gtttgagact tgctaccaga tcgctcttgc gatcaagaag gaagtcgagg 180
 atctcgaggc tgggtggtatt cagggttatcc aaatcgacga ggctgcactg agagaagggc 240
 tgccgtccgc aaggctgagc atgcatt 267

<210> 1182
 <211> 249
 <212> nucleic acid
 <213> Zea mays

<400> 1182

tgtgaagtac ggagctggca ttggccctgg tgtctacgac atccactctc ctaggattcc 60
 ctccacagag gagatcgag accgcgtcga gaagatgctc gccgtgctcg acaccaacat 120
 cctctgggtg aaccctgact gtgggtctcaa gacacgcaag tacacggagg tcaagcccgc 180
 cctgaccaac atgggtctcg ccaccaagct catccgcacc cagcttgcca gcgcgaaatg 240
 aggtcgttt 249

<210> 1183
 <211> 259
 <212> nucleic acid
 <213> Zea mays

<400> 1183

ctcacctatc atcgccgctg cgcgctctcc tccgctcccg tcgtccggaa agatggcgctc 60
 ccacattggt ggataccctc gcatgggccc caagagggng ctcaagtttg cacttgagtc 120
 tttctgggat gggaagagca ccgctgagga tctggagaag gttgctaccg acctcagggc 180
 cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca acactttctc 240
 gtactatgac caggtcctt 259

<210> 1184
 <211> 296
 <212> nucleic acid
 <213> Zea mays

<400> 1184

ctaccaaata gctcttgcaa tcaaaaagga ggttgaggat cttgangctg ctggtattca 60
 ggtgatccag atcgatgagg cagctctaag ggagggctctg ccactacgca agtcagagca 120

tgcattctac ctggactggg ctgtccactc tttcaggatc accaactgcg gagtccagga 180
caccacccag atccacaccc acatgtgcta ctccaacttc aacgacatca tccactccat 240
ccatcgacat gggatgccga tgtgatcacg atcgagaact cccggtctga ngagaa 296

<210> 1185
<211> 218
<212> nucleic acid
<213> Zea mays

<400> 1185

aggctgcact gagagaaggg ctgccgctcc gcaaggctga gcatgcattc tacttggact 60
gggctgtcca ctcttcaga atcaccaact gcgagatcca ggacaccacc cagatccaca 120
cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac atggacgcgg 180
acgtgatcac catcgagaac tcgcgggtccg acgagaag 218

<210> 1186
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 1186

ctctcacct atcatcgccg ctgcgccgtc tctccgctc ccgtcgtccg gaaagatggc 60
gtccacatt gttggatacc ctgcgatggg cccaagagg gagtcaagt ttgcacttga 120
gtttttctgg gatgggaaga gcaccgctga ggatctggag aagggttgta ccgacctcag 180
ggccagtatc tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt 240
ctcgtactat gaccaggtcc t 261

<210> 1187
<211> 282
<212> nucleic acid
<213> Zea mays

<400> 1187

gcgtcccaca ttgttgata ccctcgcatg ggccccaaga gggagctcaa gtttgactt 60
gagtctttct gggatgggaa gagcaccgct gaggatctgg agaagggtgc taccgacctc 120
agggccagta tctggaagca gatggctgat gctgggatca agtacatccc cagcaacact 180

ttctcgtact atgaccaggt ccttgacaac atgccatgct cggcgtgtcc ccgaacgtta 240
ctcatggatn gaggagagat ggattgacaa ctattctcca tg 282

<210> 1188
<211> 298
<212> nucleic acid
<213> Zea mays

<400> 1188

gtgcgtaag cctccgatca tctacgggtga cgtgagccgc cccaacccca tgacnntttt 60
ctggtcgaag aggcacagag catgacgtct cgcccaatga aggggatgct gactggccca 120
gtcacaatcc tcaatggtcc ttctgtccga atgaccagcc gaggtttgag acttgctacc 180
agatcgctct tgcgatcaag aaggaagtcg aggatcttga ggctgggtggg attcaggtta 240
tccaaatoga cgaggctgcc actgagagaa gggctgccgt ccgcaaggct gagcatca 298

<210> 1189
<211> 228
<212> nucleic acid
<213> Zea mays

<400> 1189

cccagatcca caccacatg tgctactcca acttcaacga catcatccac tncgatcatc 60
gacatggacg cggacgtgat caccatcgag aactcgcggg ccgacgagaa gctgctctcc 120
gtgttccgtg aggggtgtcaa gtacggcgcg ggcacggcc ccggtgtcta cgacatccac 180
tccccagga tcccgtcggc cgaggagatc gccgaccga tcgacaag 228

<210> 1190
<211> 280
<212> nucleic acid
<213> Zea mays

<400> 1190

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caaagaagat caccgaggac gaatacatca gtgccatcaa ggaagaaatc agcaaggncg 120
tcaagatcca agaggagctt gacattgatg tgcttgtgca tggagagcca gagagaaatg 180
acatggttga gtacttcggg gagcaattat ctggtttgcg ttcactgcca acggatgggt 240

gcaatcctat ggatcacgct tgtgtgaagc caccattat 280

<210> 1191
<211> 290
<212> nucleic acid
<213> Zea mays

<400> 1191

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ttgtgaacga gactaagctt gacagcgaga ttaagtctng gcttgctttt gctgcccaga 120
aggttggtga gggtgatgct cttgctaagg cattggctgg tcaaaaggat gaggcttact 180
tcgcagcaaa tgctgctgct caggcctcga ggaaatcctc acccgtgtg accatgaaga 240
agtccagaag gctgctgctg ctctcaaggg tctgaccacc gccgtgggac 290

<210> 1192
<211> 243
<212> nucleic acid
<213> Zea mays

<400> 1192

gaccagccga gggttgagac ttgctaccag atcgctcttg cgatcaagaa ggaagtcgag 60
gatcttgagg ctgggtggtat tcagggttatc caaatcgacg aggctgcact gagagaaggg 120
ctgccgctcc gcaaggctga gcatgcattc tacttggtact gggctgtcca ctcttcaga 180
attaccaact gcgagatcca ggacaccacc cagatccaca cccacatgtg ctactccaat 240
tca 243

<210> 1193
<211> 223
<212> nucleic acid
<213> Zea mays

<400> 1193

atggacgcag atgtgatcac catcgagaac tcccggtccg acgagaagct actgtccgtg 60
ttccgcgagg gtgtgaagta cggagctgga atcggtccag gtgtctacga catccactcg 120
cccaggatcc cctccaccga ggagatcgcg gaccgctca acaagatgct cgccgtgctc 180
gacaccaaca tcctctgggt gaaccctgac tgcggtctca aga 223

<210> 1194
 <211> 246
 <212> nucleic acid
 <213> Zea mays

 <400> 1194

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 tgagacttgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc tcgaggctgg 120
 tggatattcag gttatccaaa tcgacgaggc tgactgaga gaagggtgc cgctccgcaa 180
 ggctgagcat gcattctact tggactgggc tgtccactcc ttnagantca ccacngggag 240
 atccng 246

<210> 1195
 <211> 232
 <212> nucleic acid
 <213> Zea mays

 <400> 1195

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 caaggctgan catgcattct acttgactn ggctgtccac tccttcagaa ttaccaactg 120
 cgagatccag gacaccaccc agatccacac ccacatgtgc tactccaact tcaacgacat 180
 catccactcg atcatcgaca tggacgcgga cgtgatcacc attgagaact cg 232

<210> 1196
 <211> 248
 <212> nucleic acid
 <213> Zea mays

 <400> 1196

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 gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg ttttgcggtc 120
 actgccaatg gctgggtgca gtcttatggg tcaagggtgcg tcaagcctcc gatcatctac 180
 ggtgacgtga gccgccccaa ccccatgacc gttttctggt cgaagaggcg cagagcatga 240
 cgtctcgc 248

<210> 1197

<211> 300
 <212> nucleic acid
 <213> Zea mays
 <400> 1197
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 cttgaggctg ttgttgggaa ggacaagctt gtcgtatcaa cttcctgctc gctcatgcac 120
 accgctgtgg atcttgtgaa cgagactaag cttgacagcg agattaagtc ttggcttgct 180
 tttgctgccc agaaggttgt tgaggttgat gctcttgcta aggcattggc tggcmetaag 240
 gatgaggcta cttcgcagca aatgctgctg ctcaggcctc gaggaatcc tcaccccgctg 300

<210> 1198
 <211> 233
 <212> nucleic acid
 <213> Zea mays
 <400> 1198
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 tgcattctac ctggactggg ctgtccactc tttcaggatc accaactgcg gagtccagga 120
 caccacccag atccacaccc acatgtgcta ctccaacttc aacgacatca tccactccat 180
 catcgacatg gatgccgatg tgatcacgat cgagaatccc ggtctgacga gaa 233

<210> 1199
 <211> 256
 <212> nucleic acid
 <213> Zea mays
 <400> 1199
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 gtcgaggatt cttggaggct ggtggtattc aggttatcca aatcgacgag gctgcactga 120
 gagaagggct gccgctccgc aaggctgagc atgcattcta cttggactgg gctgtccact 180
 ccttcagaat taccaactgc gagatccagg acaccaccca gatccacacc cacatgtgct 240
 actccaactt caacga 256

<210> 1200
 <211> 297
 <212> nucleic acid

<213> Zea mays

<400> 1200

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aggattccct ccaccgagga gatcgagac cgctcgaga agatgctgc cgtgctcgac 120
accaacatcc tctgggtgaa cctgactgt ggtctcaaga cacgcaagta cacggaggtc 180
aagccccccc tgaccaacat ggtctcgcc accaagctca tccgcaccca gcttgccagc 240
gcgaaatgag tcgtttgata gctccatggt ctgatagcgc ggaatgagcc agtttgt 297

<210> 1201

<211> 260

<212> nucleic acid

<213> Zea mays

<400> 1201

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acattgttgg ataccctcgc atgggccccca agaggagct caagtttgca cttgagtctt 120
tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggccca 180
gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac ancttctcgt 240
actatgacca ggtcttgaca 260

<210> 1202

<211> 261

<212> nucleic acid

<213> Zea mays

<400> 1202

gttccgcgag ggcgtcaagt acggcgcggg catcgcccc ggtgtctacg acatccactc 60
ccccaggatc ccgtcggctg aggagatcgc cgaccgcatc gacaagatgc tggcgggtgct 120
ggacaccaac atcctctggg tgaacccccga ctgcggcctc aagacncgca agacccacgg 180
aggtcaagcc cncctgaen aacatggtct ccgccgctaa gctcatncgc acccagntcg 240
ccagcgccaa gtgagcactt t 261

<210> 1203

<211> 246

<212> nucleic acid

<213> Zea mays

<400> 1203

cccttgccaa ggctttggca ggccaaaagg atgaggtcta ctttgcagcc aatgctgctg 60
ctnnggcctc aaggagatca tcgcccaggg tgacaaaacga ggaggtccag aaggctgcag 120
ctgctttgag gggatctgac caccgccgtt ctaccactgt ttctgctaga ttggatgctc 180
agcagaaaaa gctcaacctt cctgtccttc ccacaaccac aattggttca ttccctcaga 240
ctgtgg 246

<210> 1204

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 1204

gggacntggt ccagtgggta ncgcggacgt gatcaccatt ganaactcgc ggtccgaagn 60
gaagctgctc tccgtgttcc gcnagggcgt caagtacggc gcgggcatcg gccccggtgt 120
ctacgacatc cactccccca ggatcccgtc ggctgaggat atcgccgacc gcatcgacaa 180
gatgctggcg gtgetggaca ccaacatcct ctgggtgaac cccgactgcg gcctcaagac 240
ccncaagtac acggaggctc agcccc 266

<210> 1205

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 1205

ccccaacccc atgaccgttt tcnggtcgaa gacggcacag agcatgacgt ctgcccacat 60
gaaggggatg ctgactggcc cagtcacaat cctcaactgg tccttcgtcc ggaatgacca 120
gcogangttt gagacttgct accagatcgc tcttgcatc aagaaggaag tcgaggatct 180
tgagtntggt ggtattcagg ttatccaaat cgacgaggct gcactgagan aagggcnnnn 240
cgctccgcaa ggctgagcat gcattctaata tgganggntc atccttcaga ttacactgng 300
ag 302

<210> 1206

<211> 319
 <212> nucleic acid
 <213> Zea mays

 <400> 1206

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 caaccccatg accgttttct ggtcgaagac ggcgagagc atgacgtctc gcccaatgaa 120
 ggggatgctg actggcccag tcacaatcct caactggctc ttcgtccgga atgaccagcc 180
 gaggtttgag acttgctacc agatcgctct tgcgatcaag aaggaatcga ggactcgagg 240
 ctggtggtat cagttaccaa ncgagagtgc ntgagagaag gtgcgtcgca gntgacnnat 300
 ctattgatgg tgtcatctc 319

<210> 1207
 <211> 299
 <212> nucleic acid
 <213> Zea mays

 <400> 1207

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 tgggtgttgat gatggaagca acatctgggc tgatgatctt gctacatctc tcagcantct 120
 ccagtctctt gaggtgttg ttgggaagga caagcttgct gtatcaactt cctgctcgct 180
 catgcacacc gctgtggatc ttgtgaacga gactaagctt gacagcgaga ttaagtcttg 240
 gcttgctttt gctgcccaga angttgttga gttgatgtct tgctaaaggc attgctggt 299

<210> 1208
 <211> 315
 <212> nucleic acid
 <213> Zea mays

 <400> 1208

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 antncttggc ctgntttgct gcccagaagg ttgttgaggt tgatgctctt gctaaggcat 120
 tggtggtca aaaggatgag gcttacttcg cagcaaagtc tgcngtcagg cctcgaggaa 180
 atcctcacc cgtgtgacca atgaagaagt ccagaaggct gntgctgctc tcaagggctc 240
 tgaccaccgc cgtgggacca acgttagtgc tagatggatg cccagcngtn agnagcttaa 300

cctccccanc cttcc

315

<210> 1209
<211> 421
<212> nucleic acid
<213> Zea mays

<400> 1209

cccactcttt cgtgggtnag gacaagcttg tcntatcaag ttcctggggg gctcatgcac 60
accgntgtgg atcttgtgaa cgagactaat cttgacagcg agattaagtc ttggcttgct 120
tttgctgccc agaaggttgt tgaggttgat gctcttgcta aggcatggc tggcctaaaag 180
gatgaggctt acttcgcagc aaatgctgct gctcaggcct cgaggaaatc tcaccccgtn 240
tgaccaatga anaantccan aaggctgctg ctgctctcaa gggctctgac caccggcgtn 300
ggaccaacgt tagtgctaga ttggatgccc acaaangaaa cttaacctcc cattcttcca 360
ccacttcaat tggttcgtnn ccacaaaccc gttgagctca agaagggtccg ccgtgagtac 420
a 421

<210> 1210
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 1210

catcggtccc ggtgtctacg acatccactc cccaggatc ccgtcggtccg aggagatcgc 60
cgaccgcacg gacaagatgc tggccgtgct ggacaccaac atcctctggg tgaaccccca 120
ctgcggcctc aagaccgcga agtacacgga ggtcaagccc gccctgacca acatggtctc 180
cgctgctaag ctcatcgca cccagctcgc cagcgccaag tgagcacctt tttttgcctt 240
tttcgtttcc gaggagggcg t 261

<210> 1211
<211> 239
<212> nucleic acid
<213> Zea mays

<400> 1211

gggaaggaca agcttgctgt atcaacttcc tgctcgctca tgcacaccgc tgtggatctt 60

gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgcttttgc tgcccagaag 120
gtttgtgagg ttgatgctct tgctaaggca ttggctggtc aaaaggatga ggcttacttc 180
gcagcaaatg ctgctgctca ggctcggagg aaatcctcac cccgtgtgac caatgaaga 239

<210> 1212
<211> 236
<212> nucleic acid
<213> Zea mays

<400> 1212

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ggataccctc gcatggggcc caagaggagg ctcaagtttg cacttgagtc tttctgggat 120
gggaagagca ccgctgagga tctggagaag gttgctaccg acctcagggc cagtatctgg 180
aagcagatgg ctgatgctgg gatcaagtac atcccagca acactttctc gtacta 236

<210> 1213
<211> 276
<212> nucleic acid
<213> Zea mays

<400> 1213

cccacatctc agttcacccc ggcgctgctc tcttcccccg gcgctactct cccgctccac 60
gttccaaagg aaagatggcg tcccatattg ttggataccc tcgcatgggc cccaagagag 120
agctcaagtt tgccttggag tctttctggg atgggaagag cagcgccgat gatttagaga 180
aagttgcctc tgacctcagg tctagcatct ggaagcaaat gtcagaagct gggatcaagt 240
acattcccag caacaccttc tcgtactatg accagg 276

<210> 1214
<211> 231
<212> nucleic acid
<213> Zea mays

<400> 1214

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tctggctcaa gacggcgcag agcatgacgt ctgcaccaat gaaggggatg ctgantggcc 120
cagtcacaat cctcaactgg tcttctgctc ggaatgacca gccgaggttt gaganttgcn 180

ancagatcgc tcttgcgac aagaaggaag tngaggatct cgangctagt g 231

<210> 1215
 <211> 284
 <212> nucleic acid
 <213> Zea mays

<400> 1215

ctcttgtcct cgaccttgat tctgacaaat tggctgcatt ctctgctgca tacgcagaac 60
 ttgaatctgt actttctgga ttgaatgtgc ttgttgagac ttactttgct gatgttcctg 120
 ctgagtctca caagacccta acatctctga gcagtgtgac tgcttatggg tttgatcttg 180
 tccgtggaac ccaaactctt gggcttgtca cgagtgtggt tttccctgct ggaaagtacc 240
 tctttgctgg tgttgtggat ggacgcaaca tctgggctga tgat 284

1215
 284
 nucleic acid
 Zea mays
 1216
 280
 nucleic acid
 Zea mays
 1216
 60
 120
 180
 240
 280

<210> 1216
 <211> 280
 <212> nucleic acid
 <213> Zea mays

<400> 1216

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 tgttgtggat ggacgcaaca tctgggctga tgatcttgct acatctctca gcactctcca 120
 gtctcttgag gctgttgttg ggaaggacaa gcttgtcgta tcaacttcct gctcgctcat 180
 gcacaccgct gtggatcttg tgaacgagac taagcttgac agcgagatta agtcttggct 240
 gcttttgcctg cccagaaggt tgttgagggt gatgctcttg 280

<210> 1217
 <211> 283
 <212> nucleic acid
 <213> Zea mays

<400> 1217

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 tcatggactg gaggagagat tgggtttgac acctacttct ccatggccag gggcaacgcc 120
 actgtccctg ctatggagat gaccaagtgg ttgacacca actaccactt tattgtccct 180
 gaattgggcc caaacaccaa gttctcctat gcttctcaca aggctgttaa cgaatataag 240

gaggctagggc gctcgggtggt gatacgtgcc agtattgntg gac 283

<210> 1218
 <211> 337
 <212> nucleic acid
 <213> Zea mays

<400> 1218

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 ancttgatcc tgacaaattg gctgcattct ctgctgcata cgcagaaactt gaatctgtac 180
 tttctggatt gaatgtgctt gttgagactt actttgctga tgntcctgct gantcctaca 240
 agaccctaac atctctgagc agtgtgactg cttatggttt tgatctgtcc gtngaaccca 300
 aaccttgggc ttgtcacgag tgctggntcc ctgctgg 337

<210> 1219
 <211> 446
 <212> nucleic acid
 <213> Zea mays

<400> 1219

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 agatcgcgga ccgcgtcaac aagatgctcg ccgtgctgga caccaacatc ctctgggtga 120
 accctgactg cgggtctcaag acacgcaagt acacggaggt gaagcctgcc ctgaccaaca 180
 tgggtctccgc caccaagctc atccgcaccc agctcggcag cgcgaaatga gtcagatttg 240
 gtagatccat ggtttggtga acacaaaagc cagcagggtc atttgtttcg aataatttgg 300
 gtccttactc ctgttccatg gcgttaggtt aagcctcatt ggtgttggtg gggtagccggc 360
 gttacgtgtc cgtgtttctaa agtttggaat gtgtgtttct ctttttttgt nggggtttgn 420
 gtgccttttg gtatataaac agaaat 446

<210> 1220
 <211> 228
 <212> nucleic acid
 <213> Zea mays

<400> 1220

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 tgtcaagntc caagaggagc ttgacontga tgtgcttggt catggagagc cagaaagaan 120
 tgacatgggt gagtacttcg gtgagcantt atctggtttt gcantcactg ccaacggntg 180
 ggtgcnatct ttatggatca cgttgtgtga agccaccntt atctatgg 228

<210> 1221
 <211> 232
 <212> nucleic acid
 <213> Zea mays
 <400> 1221

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 gcgagattaa gtcttggtt gcttttgctg ccagaaggt tgttgagggt gatgctcttg 120
 ctaaggcatt ggctgggtcaa aaggatgagg cttacttcgc agcaaagtct gctgctcagg 180
 cctcgaggaa atcctcacc cgtgtgacca atgaagaagt ccagaaggct gc 232

<210> 1222
 <211> 217
 <212> nucleic acid
 <213> Zea mays
 <400> 1222

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 gtctgccact gcgcaagtca gagcatgcat tctacctgga ctgggctgtc cactctttca 120
 gaatcacnaa ctgctggagtc caggacacca ccagatcca naccatattg tgctactcca 180
 acttcaacga catcatccac tccatcatcg acatgga 217

<210> 1223
 <211> 277
 <212> nucleic acid
 <213> Zea mays
 <400> 1223

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 aaagcccgcc aagggtgtgg agaagggtt ccctcttctt tcccttctta gcagcatcct 120
 ccagctctac aaggagggtca ttgctgagtt gaaggcagct ggggcttcat ggattcagtt 180

tgatgagccc actcttgtcc tcgaccttga ttctgacaaa ttggctgcat tctctgctgc 240
 atacgcagaa cttgaatctg tattntgant tnatggg 277

<210> 1224
 <211> 206
 <212> nucleic acid
 <213> Zea mays
 <400> 1224

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 aagatccaag aggagcttga cattgatgtg cttgtgcatg gagagccaga gagaaatgac 120
 atgggttgagt acttcgggtga gcaattatct ggttttgcgt tcaactgccaa cggatgggtg 180
 caatcctatg gntcacgctg tgtgac 206

<210> 1225
 <211> 354
 <212> nucleic acid
 <213> Zea mays
 <400> 1225

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 tgcagtctta tgggtcaagg tgngtcaagc ctgccgattc atctactgna acgtgtatgc 120
 cgccaccaac nccatgaccg ttttctggtc gnnnacggcg cagagcatga cgtctcgccc 180
 aatgaatggg atgntgactn gncagtcac antcctcaac tngtctctncg tcggaatgac 240
 cagccgaggt ntganacttg ctaccagatc gctctngcga tnaagnnggn agtcgaggtc 300
 tcgacgtggn gtatcaggtg accaantcga cnagctgcat gagagagagc gcgc 354

<210> 1226
 <211> 228
 <212> nucleic acid
 <213> Zea mays
 <400> 1226

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 cgtcaagcct ccgatcatct acggtgacgt gagccgcccc aaccccatga ccgttttctg 120
 gtogaagagg cgcagagcat gacgtctcgc ccaatgaagg ggatgctgac tggcccagtc 180

acaatcctca actggtcctt cgtccggaat naccagccga ggttttat 228

<210> 1227
<211> 260
<212> nucleic acid
<213> Zea mays

<400> 1227

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cttgcaatca aaaaggaggt tgaggatctt gangctgctg gtaantcagg tgatccagat 120
cgatgaggca gctctaaggg agggctctgcc actacgcaag tcagagcatg cattctacct 180
ggactgggct gtccactctt tcaggatcac caactgggga gtccaggaca ccaccagat 240
ccacacccac atgtgctact 260

<210> 1228
<211> 276
<212> nucleic acid
<213> Zea mays

<400> 1228

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tcatggactg gaggagagat tgggtttgac acctacttct ccattggccag gggcaacgcc 120
actgtccctg ctatggagat gaccaagtgg ttgacacca actaccactt tattgtccct 180
gaattgggcc caaacaccaa gttctoctat gcttctcana aggctgttaa ngaatataag 240
aggctaaggc gctcgggtgtt aanaccggcc aatant 276

<210> 1229
<211> 256
<212> nucleic acid
<213> Zea mays

<400> 1229

gtacttggtg gaccagtctc gtacctgttg ctctcaaagc ctgccaaggg tgtggagaag 60
ggattccctc ttctttccct tcttagcagc atcctcccag tctacaagga ggatcattgct 120
gagttgaagg cagctggggc ttcattggatt cagtttgatg agcccactct tgtcctcgac 180
cttgattctg acaaattggc tgcattctct gctgcatacg cagaacttga atctgtactt 240

tctggattga atgtgc

256

<210> 1230
<211> 257
<212> nucleic acid
<213> Zea mays

<400> 1230

ggtgcgtcaa gcctccgata atctacgggtg acgtgagccg ccccaaacc catgaccgtt 60
ttctggctga agaggcacag agcatgacgt ctgcaccaat gaaggggatg ctgactggcc 120
cagtcacaat cctcaactgg tccttcgtcc ggaatgacca gccgagggtt gagacttgct 180
accagatcgc tcttgcgata aagaaggaag tcgaggatct tgaggctggt ggtattcagg 240
ttatccaaat cgacgag 257

<210> 1231
<211> 344
<212> nucleic acid
<213> Zea mays

<400> 1231

gggattccct cttctttccc ttcttagcag cactctccca gtctacaagg aggtcattgc 60
tgagttgaag gcagctgggg cttcatggat tcagtttgat gagccactc ttgtcctcga 120
ccttgattct gacaaattgg ctgcattctc tgctgcatac gcagaacttg aatctgtact 180
ttctggattg aatgtgcttg ttgagactta ctttgctgat gttcctgctg agtcctacaa 240
gaccctaaca tccctgagca gtgtgactgc ttatggtttg acttgctccgt ggaacccaaa 300
ctcttggggt gtcacgagtg ctggtttccc tgctggaagt actc 344

<210> 1232
<211> 278
<212> nucleic acid
<213> Zea mays

<400> 1232

gtccacatt gttggatacc ctgcgatgg cccaagagg gagcnccaag tttgcacttg 60
agtctttctg ggatgggaag annaccgctg aggatctgga gaaggttgct accgacctca 120
gggccagtat ctggaagcag atggnngatg ctggggatca agtacatccc cagcaacact 180

ttctcgtact atgaccaggt ccttgacacc actgccatgc tcggcgctgt ccccgaaact 240
tactcatgga ctggaggaga gattggattt gacactac 278

<210> 1233
<211> 249
<212> nucleic acid
<213> Zea mays

<400> 1233

cttctcgtc gctcatgcac accgctgtgg atcttgtgaa cgagactaag cttgacagcg 60
agattaagtc ttggcttgct tttgctgccc agaaggttgt tgaggttgat gctcttgcta 120
aggcattggc tgggtcaaaag gatgaggctt actcgcagca aatgctgcag ctcaggcctc 180
gaggaaatcc tcaccccgctg tgaccaatga agaagtcag aaggctgctg ctgctctcaa 240
gggctctga 249

<210> 1234
<211> 295
<212> nucleic acid
<213> Zea mays

<400> 1234

accaccaca tctcagttca nccggcgctc gtctcttcc cccgngcta ctctcccgct 60
ccacgttcca aaggantgct ggcgtcccat attgttggat accntcgcac gggccccaag 120
agagagctca agtttgcctt ggagtcttcc tgggatggga agagcagcgc cgatgattta 180
gagaaagttg cctctgacct caggtctagc atctggacng ccatgtcag angtgggnt 240
cnagtacatt cccagcaaca ccttctcgta ctatgaccag gtccttgata ccagg 295

<210> 1235
<211> 267
<212> nucleic acid
<213> Zea mays

<400> 1235

aacccaaact cttgggcttg tccacgagtg ctggtttccc tgctggaaaag tacctctttg 60
ctgggtgttg ggatggacgc aacatctggg ctgatgatct tgctacatct ctcagcactc 120
tccagtctct tgaggctgtt gttgggaagg acaagcttgt cgtatcaact tctgtctgc 180

tnntgcacac cgctgtggat cttgtgaacg agactaagct tgacagcgag attaatgtctg 240
gctgcttttg ctgcccagaa ggttggt 267

<210> 1236
<211> 292
<212> nucleic acid
<213> Zea mays

<400> 1236

attggccctg gtgtctacga catccactct cctaggattc cctccaccga ggagatcgca 60
gaccgcgtcg agaagatgct cgccgtgctg gacaccaaca tcctctgggt gaaccctgac 120
tgtggtctca agacacgcaa gtacacggag gtcaagcccg cctgaccaa catggtctcg 180
gccancaagc tcatccgcac ccagcttgcc aacgcgaaat gagtcgtttg atagctccat 240
ggtttgatag cgcggaatga gccagttggt ttgaataatt tgggtgntac cc 292

<210> 1237
<211> 217
<212> nucleic acid
<213> Zea mays

<400> 1237

ccaagaanan cttgacattg atgtgcttgt gcatggcgag cctnagagaa acgatatggt 60
tgagtacttc ngtgagcagc tctccggttt tgcgttcaact gccaatggct ggggtgcagtc 120
ttatgggtca aggtncgtca ngcctccgnt catctacggt gacgtgagcc gcccacccc 180
catgaccgtn ttctggtcgn agactgcgca gagcatg 217

<210> 1238
<211> 456
<212> nucleic acid
<213> Zea mays

<400> 1238

gtacggcgcg ggcacgnc cgggtgtctn tnntatccac tccccagga tcccgctggg 60
ngaggntatc gccgaccgca tcgacaagat gctggcggtg ctggacacca acatcctctg 120
gggtgaacccc gactgcggcc tcaagacccg caagtacacg gaggtcaagc ccgcctgac 180
caacatggtc tccgccccta agtcatccn caccagctc gccagcgcca agtgagcacc 240

ttttttttgc ctttttctgtt tccgaggagg gcgtcgtcna ntgccaattt ggttncaata 300
 aacaggggctn tgcccgccgt tctgttgtac tccgtctgtg gttagggttag tagttatctt 360
 gatctcgcnn ncacgccggt acctntttac tactctctgt ttggtgggtt totgaagcaa 420
 gttgcccntt tacttgtatn gtaaaaaccc aatngt 456

<210> 1239
 <211> 289
 <212> nucleic acid
 <213> Zea mays

<400> 1239

cgccgctgcg ccgtctcttc ngtnccgtc gtccggaaag atggcgtncc acattgttgg 60
 ataccctcnc atgggccccca agagggagct caagtttgca cttgagtctt tctgggatgg 120
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180
 gcagatggct gatgctggga tcaagtacat cccagcaac atttctcgta ctatgaccag 240
 tcctgacacc atgccatgtc ggcgctgtcc ccgaaggtae tcatggatg 289

<210> 1240
 <211> 274
 <212> nucleic acid
 <213> Zea mays

<400> 1240

ggaggctaag gcgctcgggtg ttgataccgt gccagtactt gttggaccan tctcgtacct 60
 gttgctctca aagcctgccca aggggtgtgga gaagggattc cctcttcttt cccttcttag 120
 cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggcttcatg 180
 gattcagttt gatgagccca ctcttgctct ngacctgaat tcggnacaaa tnggctgcat 240
 tctcngcngc atangcngaa nttgaatctg tact 274

<210> 1241
 <211> 268
 <212> nucleic acid
 <213> Zea mays

<400> 1241

atcgngctg cgccgtctcc tccgtccccg tcgtccggaa agatggcgtc ccacattgtt 60

ggataccctc gcatgggncc cnaganggag ctcaagtttg cacttgagtc tttctgggat 120
 ggggnagagca acgctgagga tctggagaag gttgctaccg acntcagggc cagtatccgg 180
 aagcanatgg cngatnctgg gatcaagtac atccccagca anacnttctc gnantatgac 240
 caggtcttga naccatgccca tggtctcg 268

<210> 1242
 <211> 260
 <212> nucleic acid
 <213> Zea mays

<400> 1242

acgacatcca ctccccagg atcccgctcg ctgaggagat cgccgaccgc atcgacaaga 60
 tgctggcggt gctggacacc aacatcctct ggggtgaacc cgactgcggc ctcaagacct 120
 gcaagtacac ggaggtcaag cccgcctga ccaacatggt ctccgccgct aagctcatcc 180
 gcaccagct cgccagcgcc aagtgagcac cttttttttg cttttttcgt ttccgaggag 240
 ggcgtcgtcg atgccaattt 260

<210> 1243
 <211> 280
 <212> nucleic acid
 <213> Zea mays

<400> 1243

ctcgcggtcc gacgagaagc tgctctccgt gtnccgcgag ggcgnccaag tacggcgcg 60
 gcatcgcccc cgggtgtctac gacatccact cccccaggat cccgtcggct gaggagatcg 120
 ccgaccgcat cgacaagatg ctggngtgct ggacaccaac atcctctggg tgaaccccca 180
 ctgcggcctc aagaccgca agtacacgga ggtcaagccc gccctgacca ncatggtctc 240
 ngccggtaag ctnatccgca ccagctcgc naggccaagt 280

<210> 1244
 <211> 305
 <212> nucleic acid
 <213> Zea mays

<400> 1244

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aacttgaatc tgtactttct ggattgaatg tgcttggtga gaacttacttt gctgatgttc 120
 ctgctgagtc ctacaagacc ctaacanctc tgagcagtg gactgctnat ggttttgata 180
 ttgtccgtgg aacccaaact ctnggggctt gtcacgagtg ctgggtttcc ctgctggaaa 240
 gtacctcttt gctgggtgtg tggatggacn caacagctgg ggtngatgnt cttncnnacn 300
 tcgcc 305

<210> 1245
 <211> 198
 <212> nucleic acid
 <213> Zea mays

<400> 1245

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 atgangggga tgctgactgg cccagtcaca atcctcaact ggtccttcgt ccggaatgac 120
 cagccgaggt ttgagacttg ctaccagatc gctcttgca tcaagaagga agtcgaggat 180
 ctcgaggctg gtggtatt 198

<210> 1246
 <211> 297
 <212> nucleic acid
 <213> Zea mays

<400> 1246

ctcccagtct acaaggaggt cattgctgag ttgaaggcag ctggggcttc atggattcag 60
 tttgatgagc ccaactcttg cctcgacctt gattctgaca aattggctgc attctctgct 120
 gcatacgag aacttgaatc tgtacttctg gattgaatgt gctgttgaga cttactttgc 180
 tgatgttctt gctgagtcct acaagaccct aacatctctg agcagtgtga ctgcctatgg 240
 ttttgatctt gtccgtggaa cccaaactct tgggcttgct acgagtgctg gtttccc 297

<210> 1247
 <211> 286
 <212> nucleic acid
 <213> Zea mays

<400> 1247

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gagaagctgc tctccgtgtt ccgcgagggc gtcaagtacg gcgcgggcat cggccccggg 120
 gtctacgaca tccactcnnc caggatccccg tcggctgagg agatcgccga ccgcacgac 180
 aagatgtgng gtgctggana cnaacatcct ctnggtgaac cccgactgog gcctcaagan 240
 ccgcaagtac acngagggtca aancggngctg accaacangt ctccgt 286

<210> 1248
 <211> 343
 <212> nucleic acid
 <213> Zea mays
 <400> 1248

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 tgacgtctca gcccaatgaa nggggnntg actggccaag tcacaatncc tcaactggtc 120
 cttcgtccgg aatgaccagc cgaggtttga gacttgctac cagatcgctc ttgcgatcaa 180
 gaaggaagtc gaggatctcg aggctgggtgg tattcagggtt atccaaatcg acgaggctgc 240
 actgagagaa gggctgccgc tccgcaaggc tgnagcatgc atcttacttg gatgggctgt 300
 tccattcctt cagaatcacc aatgogaatc caggacacca ccc 343

<210> 1249
 <211> 210
 <212> nucleic acid
 <213> Zea mays
 <400> 1249

gagacatgct accaaatagc tcttgcaatc aaaaaggagg ttgaggatct tgaggctgct 60
 ggtattcagg tgatccagat cgatgaggca gctctaaggg agggctctgcc actacgcaag 120
 tcagagcatg cattctacct ggactgggct gtccactctt tcaggatcac caactgggga 180
 gtccaggaca ccaccagat ccacaccac 210

<210> 1250
 <211> 206
 <212> nucleic acid
 <213> Zea mays
 <400> 1250

gcttttgctg ccagaagggt tgttgagggt gatgctcttg ctaaggcatt ggctgggtcaa 60

aaggatgagg cttacttcgc agcaaagtct gcagctcagg cctcgaggaa atcctcacc 120
 cgtgtgacca atgaagaagt ccagaaggct gctgctgctc tcaagggtc tgaccaccgc 180
 cgtgcgacca acgttagtgc tagntt 206

<210> 1251
 <211> 223
 <212> nucleic acid
 <213> Zea mays

<400> 1251

gaccagccga ggtttgagac ttgctaccag atcgtctcttg cgatnaagaa ggaagtcgag 60
 gatcttgagg ctggtggtat tcaggttatt caaatcgacg aggtctgact gagagaaggg 120
 ntgccgctcc gcaaggctga tcatgcattc tacttggact gggctgtcca ntcttcagn 180
 attaccaact gcgagatcca ggacaccacc cagatccaca ccc 223

<210> 1252
 <211> 186
 <212> nucleic acid
 <213> Zea mays

<400> 1252

acaaaccgtt gagctcagga gggctccgccg tgagtacaag gcaaaaaaga tctctgagga 60
 ggaatacgtc actgccatca aggaggaaat caacaaggct gttaagcttc aagaagagct 120
 tgacattgat gtgcttgtgc atggcgagcc tgagagaaac gatatggttg agtacttcgg 180
 tgagca 186

<210> 1253
 <211> 238
 <212> nucleic acid
 <213> Zea mays

<400> 1253

cnagatggcg tcccatattg ttggataccc tcgcatgggc cccaagaggg agctgcaagt 60
 ttgcnttgga gtactttctg ggatgggnag agcagcgccg aggatttgga gaangttgnc 120
 actgacctga ggtctagcat ctgnagcaa atgtcagang ctgggatcaa gtacattccc 180
 aggcaatcac cttctcgtac tacgaccagg ttcttgatac cacggccatg cttggcgc 238

<210> 1254
 <211> 277
 <212> nucleic acid
 <213> Zea mays

 <400> 1254

 gcccttgcca aggctttggc aggccaaaag gatgaggtct actttgcagc caatgctgct 60
 gcttcaggcc tcaaggagat catcgcccag ggtgacaaac gaggaggtcc agaaggctgc 120
 agctgctttg aggggatctg accaccgccc ttctaccact gtttctgcta gattggatgc 180
 tcagcagaan aagctcaacc ttctgtctct tcccacaacc acaattgggt cattccctca 240
 gactgtggaa tcaggagggt tcgccgtgaa tacaagg 277

<210> 1255
 <211> 287
 <212> nucleic acid
 <213> Zea mays

 <400> 1255

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 gtcgtcctct tccccggcgc ctactctccc gctccacgtt ccaaaggaaa gatggcgtcc 120
 catattgttg gataccctcg catgggcccc aagagagagc tcaagtttgc cttggagtct 180
 ttctgggatg ggaagagcag cgccgatgat ttagagaaag ttgcctctga cctcaggctc 240
 agcatctgga agcaaattgc agaagctggg atcaagtaca ttcccag 287

<210> 1256
 <211> 219
 <212> nucleic acid
 <213> Zea mays

 <400> 1256

 gggcccaaac accaagttct cctatgcttc tcacaaggct gttaacgaat ataaggaggc 60
 taaggcgctc ggtgttgata ccgtgccagt acttgttgga ccagtctcgt acctgttgct 120
 ctcaaagcct gccaaagggtg tggagaaggg attccctctt ctttcccttc ttagcagcat 180
 cctcccagtc tacaaggagg tcattgctga gttgaaggc 219

<210> 1257

<211> 269
 <212> nucleic acid
 <213> Zea mays

 <400> 1257

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 gaagtcgagg atctcgaggc tgggtgtatt caggttatcc aaatcgacga ggctgcactg 120
 agagaagggc tgccgctccg caaggctgag catgcattct acttggaactg ggctgtccac 180
 tccttcagaa tcaccaactg cggagatcca ggacaccacc catatccaaa cccanatgtg 240
 ctattcnanc ttnaacgaca tcatccact 269

<210> 1258
 <211> 213
 <212> nucleic acid
 <213> Zea mays

 <400> 1258

 caacttcctg ctcgctcatg cacaccgctg tggatcttgt gaacgagact aagcttgaca 60
 gcgagattaa gtcttggtt gcttttgcctg cccagaaggt tgttgagggt gatgctottg 120
 ctaaggcatt ggctgggtcaa aaggatgagg cttacttcgc agcaaagtct gctgctcagg 180
 cctcgaggaa atcctcacc cgtgtgacca atg 213

<210> 1259
 <211> 269
 <212> nucleic acid
 <213> Zea mays

 <400> 1259

 atttggttgc cctggccggt acnagtccaa gaattcccgg gtcctcacct atcatcgccg 60
 ctgcgcgctc tctccgctc ccgtcgctccg gaaagatggc gtcccacatt gttggatacc 120
 ctgcgatggg cccaagagg gagctcaagt ttgcacttga gtctttcttg gatgggaaga 180
 gcaccgctga ggatctggag aagggtgcta ccgacctcag ggccagtatc tggaagcaga 240
 tggctgatgc tgggatcaag tacatcccc 269

<210> 1260
 <211> 266
 <212> nucleic acid

<213> Zea mays

<400> 1260

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cttcgtccgg aatgaccagc cgagggttga gacttgctac cagatcgctc ttgcgatcaa 120
gaaggaagtt cgaggatctt naggtcgttg gtattcaggt tatccaaatc cgacgaggct 180
gcactgagna gaagggtgc cgctccgcaa ggccgagcat gccatctact tgnactgggc 240
tgtccactcc ttcagattac caactg 266

<210> 1261

<211> 443

<212> nucleic acid

<213> Zea mays

<400> 1261

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gaggagatcg ccgaccgcat cgacaagatg ctggcggtgc tggacaccaa catcctctgg 120
gtgaaccccg actgcggcct caagaccgc aagtacacgg aggtcaagcc cgcctgacca 180
acatggtctc cgccgctaag ctcatccgca cccagctcg cagcgccaag tgagcacott 240
atntttgcnt ntttngtttc cgaggagggc gtngtcgatg ccaattngng ttcaataaac 300
agggctgtgc ccgcccgttc tgtngactnc gctgtgggtt aagttagtan tttcttngat 360
ctcgnccca ccggttacct ggtttactac tctctgtttg gtggttcttg aagcaagtta 420
cnnctgtac ttggatagaa aaa 443

<210> 1262

<211> 258

<212> nucleic acid

<213> Zea mays

<400> 1262

ggcagctggg gcttcatgga ttcagtttga tgagccact cttgtcctcg accttgattc 60
tgacaaattg gctgcattct ctgctgcata cgcagaactt gaatctgtac tttctggatt 120
gaatgtgctt gttgagactt actttgctga tgttctgct gagtcctaca agaccctaac 180
atctctgagc agtgtgactg cttatggttt tgatcttgct cgtggaacct aaactcttgg 240

gcttgtcacg agtgctgg 258

<210> 1263
 <211> 275
 <212> nucleic acid
 <213> Zea mays

<400> 1263

ggcagctggg gcttcatgga ttcagtttga tgagcccact cttgtcctcg accttgattc 60
 tgacaaattg gctgcattct ctgctgcata cgcagaactt gaatctgtac tttctggatt 120
 gaatgtgctt gttgagactt actttgctga tgttcctgct gagtcctaca agaccetaac 180
 atctctgagc agtggtgactg cttatggttt tgatcttgct cgtggaaccc aaactcttgg 240
 gcttgtcacg agtgctgggt tccctgctgg aaagt 275

gcttgtcacg agtgctgggt tccctgctgg aaagt

<210> 1264
 <211> 253
 <212> nucleic acid
 <213> Zea mays

<400> 1264

ctacgacatc cactctccta ggattccctc cacagaggag atcncagacc gcgtcgagaa 60
 gatgctcgcc gtgctcgaca ccaacatcct ctgggtgaac cctgactgtg gtctcaagac 120
 acgcaagtac acggagggtca agcccgccct gaccaacatg gtctcggcca ccaagctcgc 180
 ncgcacccag cttgccagcg cgaaatgagg tcgtttgata gctccatggt ctgatagcgc 240
 ggaatgagcc agt 253

<210> 1265
 <211> 211
 <212> nucleic acid
 <213> Zea mays

<400> 1265

ggctgctgct ctcaagggt ctgaccaccg ccgtgggacc aacgttagtg ctagattgga 60
 tgcccagcag aagaagctta acctcccat ccttcccacc actacaatcg gttcgttccc 120
 acaaaccgtt gagctcagga gggccgccc tgagtacaag gcaaaaaaga tctctgagga 180
 agaatacgtc actggccatc naggaggaaa t 211

<210> 1266
 <211> 297
 <212> nucleic acid
 <213> Zea mays

 <400> 1266

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 gaccttgatt ctgacaaatt ggctgcattc tctgctgcat acgcagaact tgaatctgta 120
 ctttctggat tgaatgtgct gttgagactt actttgotga tgttcttgct gagtcttaca 180
 agaccctaac atctctgagc agtgtgactg cctatgggtt tgatcttgct cgtggaaccc 240
 aaactcttgg gcttgtcacg agtgtggtt tccctgctgg aaagtactct tgctggt 297

1266
 297
 nucleic acid
 Zea mays
 1267
 267
 nucleic acid
 Zea mays
 1267
 60
 120
 180
 240
 267

<210> 1267
 <211> 267
 <212> nucleic acid
 <213> Zea mays

 <400> 1267

 atccctgagc agtgtgactg cttatgggtt tgatcttgct cgtggaaccc aaactcttgg 60
 gcttgtcacg agtgtggtt tccctgctgg aaagtacctc tttgctggtg ttgtggatgg 120
 acgcaacatc tgggctgatg atcttgctac atctctcagc actctccagt ctcttgaggc 180
 tgttgttggg aaggacaagc ttgtcgtatc aacttctctg tctctcatgc acaccgctgt 240
 ggatcttctg aacgagacta agcttga 267

<210> 1268
 <211> 224
 <212> nucleic acid
 <213> Zea mays

 <400> 1268

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 tacnctcgca tggnccccaa nagggagctc aagtttgac ttnagtcttt cnggggatngg 120
 aagancaacng ctgaggatcn cgagaaggtn gcnaccgacc tcagggccag tatctggaag 180
 cagatngctg annctgggat caantacatn cccagcaaca cttt 224

<210> 1269

<211> 212
<212> nucleic acid
<213> Zea mays

<400> 1269

nnnnnnnnnn nnnnnngaca tggacgcgga cgtgatcacc atcgagaact cgcggtccga 60
cgagaagctg ctctccgtgt tccgtgaggg tgtcaagtac ggcgcgggca tcggccccgg 120
tgtctacgac atccaactccc ccaggatccc gtgggcccag gagatcgccg accgcatcga 180
caagatgctg gccgtgctgg acaccaacat ca 212

<210> 1270
<211> 292
<212> nucleic acid
<213> Zea mays

<400> 1270

cgagangctg ctctccgtgt tcngecnnggg cgtcaagtac ggngcgggcn tcggcctcgg 60
tgtctacgac atcnntccc ccaggntncc gtcnctgng gagatcgccg accgcatcga 120
caagatgctg gcggtgctgg acaccaacat cctctgggtg aaccccgact cggcctcaag 180
acccgcaagt acacggnggt caagcccgcc ctgaccaaca tggntcgcg gctnagtcac 240
ccgaccagc tcgnnagccg cgangtgagc acttttttgc cttttcgttc cg 292

<210> 1271
<211> 287
<212> nucleic acid
<213> Zea mays

<400> 1271

angcaacatc tgggctgatg atcttgctac atctctcagc actctccagt ctcttgaggc 60
tggtgttggg aaggacaagc ttgtcgtatc aacttcctgc tcgtcatgc acaccgctgt 120
ggatcttgtg aacgagacta agcttgacag cgagattaag tottggttg cttttgctgc 180
ccagaagggt gttgaggtga tgctcntgct aaggcattgg ctgggtcaaaa ggatgaggnt 240
acttcgcagc aaatgcngca gctcaggncn cgaggaaatc ctcaccc 287

<210> 1272
<211> 265
<212> nucleic acid

<213> Zea mays
 <400> 1272
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 ctccacgttc caaaggaaag atggcgtaen catattgntg gataccctcg catgggcccc 120
 aagagagagc tcaagtttgc cttggagtct ttctgggatg ggaagagcag cgccgatgat 180
 ttagagaaaag ttgcctctga cctcaggtct agcatctgga agcanatgtc agaagctggg 240
 atcaagtaca ttcccagcaa cacct 265

<210> 1273
 <211> 223
 <212> nucleic acid
 <213> Zea mays
 <400> 1273
 tagcagcatc ctcccagtct acaaggaggt cattgctgag ttgaaggcag ctggggcttc 60
 atggattcag tttgatgagc ccactcttgt cctcgacctt gattctgaca aattggctgc 120
 attctctgct gcatacgcag aacttgaatc tgtactttct ggattgaatg tgcttggtga 180
 gacttacttt gctgatgttc ctgctgagtc ctacaagacc cta 223

<210> 1274
 <211> 261
 <212> nucleic acid
 <213> Zea mays
 <400> 1274
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 gtatctggaa gcagatggct gangctggga tcaagtacat cccagcaac actttctcgt 120
 actatgacca ggtccttgac accactgcca tgctcggcgc tgtccccgaa cgttactaca 180
 tggactggag gagagattgg atttgacacc tacttctcca tggccagggg caagccactg 240
 ttctctgctat ggagatgacc a 261

<210> 1275
 <211> 283
 <212> nucleic acid
 <213> Zea mays

<400> 1275

tggtgttgat gatggacgca acatctgggc tgatgatctt gctacatctc tcagcantnt 60
ccaggctnnt gagngctggt gttgggaatg acaagottgt cgtatcaact tctgctcgc 120
tcatgcacac cgctgtggat cttgtgaacg agactaagct tgacagcgag attaagtctt 180
ggcttgcttt tgctgccag aaggttggtt aggttgatgc nctgctaagg catggctggt 240
caaaaggatg aggettactt cgcagcaaatt gctgcagctc agg 283

<210> 1276

<211> 185

<212> nucleic acid

<213> Zea mays

<400> 1276

gcgtcccaca ttgttgata cconacgat gggccccaag agggagctac aagtttgac 60
ttgagtcttn ctgggatggg aagagcaccg ctgaggatct ggagaagggt gctaccgacc 120
tcagggccag tatctggnag cagatggctg atgctgggat caagtacatc cccagcaaca 180
ctttc 185

<210> 1277

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1277

tcagtttgat gagccactc ttgtcctcga cttgattct gacanattgg ctgcaanctc 60
tgctgcatac gcagaacttg aatctgtact ttctggattg aatgtgcttg ttgagactta 120
ctttgctgat gttcctgctg agtcctacaa gaccctaaca tccctgagca gtgtgactgc 180
ttatggtttt gatcttgctc gtggaacca aactcttggg cttgtcacga gtgctgggtt 240
ccctgctgga aagtactctt tgctggtggt gtggatga 278

<210> 1278

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1278

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acaccaacat cctctgggtg aaccccgact gcggcctcaa gaccgcgaag tacacggagg 120
tcaagcccg cctgacaaaa atggtccccg ctganaagct cattcgcacc cagctcgcca 180
gcgccaagtg agcacctttt tttgcctttt tcgtttccga ggagggcgtc gtcgatgcc 240
tttgtttcca ataaacaggg cccccccct gtgcgc 276

<210> 1279
<211> 267
<212> nucleic acid
<213> Zea mays

<400> 1279

ggggcttcat ggattcagtt tgatgagccc actcttgtcc tcgacctga ttctgacaaa 60
ttggctgcat tctctgtgc atacgcagaa cttgaatctg tactttctgg attgaatgtg 120
cttggtgaga cttactttgc tgatgttccg gctgagtcct acaagaccct aacatccctg 180
agcagtgtga ctgcttatgg ttttgatctt gtccgtggaa cccaaactct tgggcttgtc 240
acgagtgtg gtttccctgc tggaaag 267

<210> 1280
<211> 271
<212> nucleic acid
<213> Zea mays

<400> 1280

ctccatgacc gttttctggt cgaagacggc gcagagcatg acgtctcgcc caatgaaggg 60
gatgctgact ggcccagtca caatcctcaa ctggtcttcg tccggaatga ccagccgagg 120
tttganactt gctaccagat cgctcttgcg atcaagaagg aagtcgagga tctcgaggct 180
ggtggtatca gttatccaaa tcgacgaggc tgcagagag aagggtgccg ctccgcaagg 240
ctgagcatgc attctattgg actgggtgtc a 271

<210> 1281
<211> 188
<212> nucleic acid
<213> Zea mays

<400> 1281

gctggtttcc ctgctggaaa gtacctcttt gctggtgttg tggatggacg caacatctgg 60
gctgatgac ttgctacatc tctcagcaact ctccagtctc ttgaggctgt tgttggaag 120
gacaagcttg tcgtatcaac ttctgctcg ctcatgcaca ccgctgtgga tcttgtgaac 180
gagactaa 188

<210> 1282
<211> 284
<212> nucleic acid
<213> Zea mays
<400> 1282

gtcgaagagg ggcgagagca tgacgtctcg cccaatgaag gggatgctga ctggcncagt 60
cacaatcctc aactggtcct tcgtccggaa tgaccagccg aggtttgaga cttgctacca 120
gatcgctctt gcgatcaaga aggaagtcca ggatctcgag gctggtggta ttcaggttat 180
ccaanttcga cgaggctgca ctgagagaag ggctgccgct ccgnaggntn agnatgattc 240
naantgggan nggggcggcc nntcttnnag aatgnnaggn ggng 284

<210> 1283
<211> 194
<212> nucleic acid
<213> Zea mays
<400> 1283

cacaaccaca attggtncat tccctcagac tgtggaactc aggaggggtc gccgtgaata 60
caaggcaaaa agatcaccca ggacgatacn tcagtgccat caaggaanaa tcagcaaggt 120
tgtcaagatc caagaggagc ttgacattga tgtgcttggt catggagagc cagagagana 180
tgacatgggt gagt 194

<210> 1284
<211> 271
<212> nucleic acid
<213> Zea mays
<400> 1284

cggtgttgat accgtgccag tacttggttg accagtctcg tacctgttgc tctcaaagcc 60
cgccaagggt gtggagaagg gattccctct tctttccctt cttagcagca tctcccagt 120

ctaaaggagg tcattgctga gttgaaggca gctggggctt catggattca gtttgatgag 180
 cccatcttgt cctcgacctt gattctgaca aattggctgc attctctgct gcatacgag 240
 aacttgaatc tgtatttctg gattgaatgt g 271

<210> 1285
 <211> 209
 <212> nucleic acid
 <213> Zea mays

<400> 1285

gtgctcaag cctccgatca tctacggtga cgtgagccgc cccaaccca tgaccgtttt 60
 ctggctgaag acgggcacag agcatgacgt ctgcccgaat gaaagggang ctgactggcc 120
 cagtcacaat cctcaactgg tcttctgctc ggaatgacca gccgaggttt gagacttgta 180
 ccagatcgct cttgctcaa gaaggaagt 209

<210> 1286
 <211> 180
 <212> nucleic acid
 <213> Zea mays

<400> 1286

cgagagcat gacgtctcgc ccaatgaagg ggatgctgac tggcccagtc acaatcctca 60
 actggctcctt cgtccggaat gaccagccga ggtttgagac ttgctaccag atcgctcttg 120
 cgatcaagac ggaantcgag gntctcnagg ctngtggtat tcaggttatc caaatcgagc 180

<210> 1287
 <211> 200
 <212> nucleic acid
 <213> Zea mays

<400> 1287

cacctatcat cgcgctgag ccgtctctc cgtcccgc gtccggaag atggcgtccc 60
 acattgttgg ataccctcgc atgggccccca agaggagct caagtttgca cttgagtctt 120
 tctgggatgg gaagagcacc gctgaggatc tggagaagg tgcctaccgac ctcagggccca 180
 gtatctggaa gcagatggct 200

<210> 1288

<211> 324
 <212> nucleic acid
 <213> Zea mays

 <400> 1288

 aatggcgagc ctgacanaca cgatatggtt gantacttcg gtgagcagct ctccggtttt 60
 gcnttcactg ccaatggctg ggtgcagtct tatgggtcaa ggtgcgtcaa gcctccgatac 120
 atctacggtg acgtgagccg ccccaacccc atgacgtttt ctggtcgaag acggcgcaga 180
 gcatgacgtc tcgcncaatg aaggggatgc tgactggccc agtcacaatc ntcaactggt 240
 cttcgtcgga acgaccagcc gaggtttgag attgctacca gatcgtcttt gcgatnaaga 300
 angaatcgan gatctgaagc tggc 324

<210> 1289
 <211> 181
 <212> nucleic acid
 <213> Zea mays

 <400> 1289

 aggaaatcct caccocgtgt gnaccaatga agaagtccag aaggctgctg ctgctotcaa 60
 gggctctgac caccgccgtg cgaccaacgt tagtgctaga ttggatgccc agcagaagaa 120
 gcttaacctc cccatccttc ccaccactac aatcggttcg ttcccacaaa ccgttgagct 180
 c 181

<210> 1290
 <211> 275
 <212> nucleic acid
 <213> Zea mays

 <400> 1290

 caccgaggag atcgcagacc gcgtcgagaa gatgctcgcc gtgctcgaca ccaacatcct 60
 ctgggtgaac cctgactgtg gtctcaagac acgcaagtac acggaggtca agcccgccct 120
 gaccaacatg gtctcggcca ccaagctcat ccgcaccag cttgccagcg cgaaatgagt 180
 cgtttgatag ctccatggtc tgatagcgcg gaatgagcca gtttgttttg aataatttgg 240
 gtgttaccac ctgttccatg gtgtnagtgt taggt 275

<210> 1291

<211> 202
 <212> nucleic acid
 <213> Zea mays

 <400> 1291

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 tttgctgatg ttntctgtga gtcttacaag accctaacat ccctgagcag tgtgactgct 120
 tatggttttt atcttgtccg tggaacccaa actcttgggc ttgtcacgag tgctggtttc 180
 cctgctggaa agtacctctt tg 202

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 1400

<210> 1292
 <211> 181
 <212> nucleic acid
 <213> Zea mays

 <400> 1292

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 cactgccatc aaggaggaaa tcaacaaggt cgtaagctc caagaagagc ttgaacattg 120
 atgtgcttgt gctggcgagc ctgagagaaa cgatatggtt gagtacttcg gtgagcagct 180
 c 181

<210> 1293
 <211> 242
 <212> nucleic acid
 <213> Zea mays

 <400> 1293

 ccagagcgct actcttgac tggaggcgag attggcttga gcacctactt ctctatggcc 60
 aggggaaatg ccactgtccc tgccatggag atgaccaagt ggtttgatac aaactaccac 120
 tttattgtcc ctgaacttgg cccaagcacc aagttcacat acgctnctca caaggctggt 180
 tctgagtacn nggnggcaaa ggggtcggcn ttgatncnat tcccatgctg ttggacanng 240
 gc 242

<210> 1294
 <211> 176
 <212> nucleic acid
 <213> Zea mays

<400> 1294

acggcgcaga gcatggacgt ctgcccgaat gaaggggatg ctgactggcc cagtcacaat 60

cctcaactgg tccttcgtcc ggaatgacca gccgaggttt gagacttgct accagatngc 120

tcttgcgatg aagaaggaag tcgaggatct cgaggctggt ggtattcagg ttatcc 176

<210> 1295

<211> 228

<212> nucleic acid

<213> Zea mays

<400> 1295

aagggtgtgg agaagggatt cctctctctt tcccttctta gcagcatcct ccagttctac 60

aaggaggtca ttgctgagtt gaaggcagct ggggcttcat ggattcagtt tgatgagccc 120

actcttgctc tcgaccttga ttctgacaaa ttggctgcat tctctgctgc atacgcagaa 180

cttgaatctg tactttctgg atgaatgtgc ttgttgagac tactttgc 228

<210> 1296

<211> 312

<212> nucleic acid

<213> Zea mays

<400> 1296

ccagttcacc acccaacctc cactcccagt tcaccccgtc gtccctcgnc gccaccactc 60

ctgcgtcccc cggcgctact cccccgctcc acgggtccaag gaaagatggc gtcccatatt 120

gttgataacc ctgcgatggg cccaagagg gagctcaagt ttgccttggg gtctttctgg 180

gatgggaaga gcagcgccga ggatttgag aaagttgcca ctgacctgag gtctagcatc 240

tggaagcaaa tgtcagaagt gggatcaagt aattcccagc aatacttctc gtatacgncc 300

aggttttgat ac 312

<210> 1297

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 1297

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tacaangagg tcattgctga gttgaaggca gctggggctt tcatggattc agtttngatg 120
agcccactct tgtcctcgac cttgaatnct gaacaaattg gctgcattct ctgctggcct 180
acgcagaact ngaatctgta ctntctggat tgaatgtgct tgttgagact tacttttgctg 240
atgttcctgc tgagtcctac aagaccctaa 270

<210> 1298
<211> 161
<212> nucleic acid
<213> Zea mays
<400> 1298

gggcgcgggg catcgccccc ggtgtctacg acatccactc cccagggatc ccgtcggccg 60
aggagatcgc cgaccgcacg gacaagatgc tggccgtgct ggacaccaac atcctctggg 120
tgaaccccga ctgcggcctc aagaccgcga agtacaggag g 161

<210> 1299
<211> 213
<212> nucleic acid
<213> Zea mays
<400> 1299

tgacagcgag attaagtctt ggantgcttt tgctgnccca gaaggttntt gaggttgatg 60
ctcttgctaa ggcattggct ggtcaaaagg atgaggctta cttcgcagca aatgctgctg 120
ctcaggcctc gaggaaatcc tcaccccgctg tgaccaatga agaagtccag aaggctgctg 180
ctgtctcaag ggctctgacc accgccgtgg gac 213

<210> 1300
<211> 194
<212> nucleic acid
<213> Zea mays
<400> 1300

cacctatcat cgccgctgcg ccgtctcttc cgctcccgct gcccgaaag atggcgctccc 60
acattgttgg ataccctcgc atgggccccca agaggagct caagtttgca cttgagtctt 120
tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcaggggcca 180
gtatctggaa gaaa 194

<210> 1301
 <211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 1301

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 gtcacgagtg ctggtttccc tgctggaaag tacctctnng ctggtgttgt ggatggacgc 120
 aacatctggg ctgatgatct tgctacatct ctacagcactc tccagtctct tgaggctgtt 180
 gttgggaagg acaagcttgt cgtatcaact tctgtctgc tcatgcacac cgctgtggga 240
 gctgtggaac cgagantagc ttganaangg ggttaagtnt nggcttgntn t 291

<210> 1302
 <211> 152
 <212> nucleic acid
 <213> Zea mays

 <400> 1302

 tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 60
 gctctccggt tttgcgttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 120
 caagcctccg atcatctacg gtgacgtgag cc 152

<210> 1303
 <211> 152
 <212> nucleic acid
 <213> Zea mays

 <400> 1303

 ggngctgtcc actccttcag aattaccaac tgcgagancc aggacaccac ccagatccan 60
 ncccacatgt gctactccaa cttaacgac atcatccact cgatcatcga catggacgcg 120
 gacgtgatca ccatngagaa ctcgngggnc gg 152

<210> 1304
 <211> 468
 <212> nucleic acid
 <213> Zea mays

 <400> 1304

cttganccttg ncccangccg gnactgggta ctactntntg gtngnggttc tgag 414

<210> 1307
 <211> 163
 <212> nucleic acid
 <213> Zea mays

<400> 1307

cttcccccg cgtactctc ccgtccacg ttccaaagga aagatggcgt cccatattgt 60

tggataccct cgcattgggc ccaagagaga gctcaagttt gccttggagt ctttctggga 120

tgggaagagc agcgccgatg atttagagaa agntgnntcg gac 163

<210> 1308
 <211> 233
 <212> nucleic acid
 <213> Zea mays

<400> 1308

caccagttca ccaccacct cccactcca gttcaccccg tcgtcctcgg cgccaccact 60

cctcgtcccc cggcgctact ccccgctcc acgggtccaag gaaagatggc gtcccatatt 120

gttgataacc ctgcattggg cccaagagg gagctcaagt ttgccttga gtctttctgg 180

gatgggaaga gcagcgccga ggatttggag aaagttgcc ctgacctgag gtc 233

<210> 1309
 <211> 225
 <212> nucleic acid
 <213> Zea mays

<400> 1309

atcaccagtt caccaccac ctccactcc cagttaccc cgtcgtcctc ggcgccacca 60

ctcctcgtcc cccggcgcta ctccccgct ccacgtcca aggaaagatg gcgtcccata 120

ttgttgata ccctcgcatt ggccccaga gggagctcaa gtttgccttg gactctttct 180

gggatgggaa gagcagcgc gaggatttgg agaaagttgc cactg 225

<210> 1310
 <211> 303
 <212> nucleic acid
 <213> Zea mays

<400> 1310

atcatcacca gttcaccacc cacctcccac tcccagttca ccccgctcgtc ctcgggcgcca 60
ccactcctcg tcccccgggc ctactccccc gctccacggg ccaaggaaag atggcggtccc 120
atattgttgg ataccctcgc atgggccccca agagggagct caagtttgcc ttggagtctt 180
tctgggatgg gaagagcagc gccgaggatt ggagaaagtt gccatgacct gangttancc 240
attggaagca aatgtcagaa ntgggatcaa gtacatccca gaatacttct cgtatagaca 300
ggt 303

<210> 1311

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 1311

gtctacaagg aggtcattgc tgagttgaan gctgggnggn tgctgctaag angggaaaga 60
ngagggaatc ctttcttagc agcatcctcc cagtctacna ggnggtcatt gctgagttga 120
aggcagctgg ggcttcatgg attcagtttg atgagccac tcttgtecg gacottgatt 180
ctgacaaatt ggctgcattc tctgctgcat acgcagaact tgaatctgta ctttctggat 240
tgaatgtgct gttgagacta cttgctgatg ttctgctga gtccatacaag acc 293

<210> 1312

<211> 311

<212> nucleic acid

<213> Zea mays

<400> 1312

ccccaacccc atgaccgttt nctggctgaa gacggcacag agcatgacgt ctgcccattg 60
aaggggtgct gactggccca gtcacaatcc ncaactgggc cttcggtccg aatgaccagc 120
cgaggtttga gacttgctac cagatcgtct tgcgatcaag aaggaagtcg aggatcttga 180
ggttgtgtat tcaggttatc caaatcgaga gctgcatgag agaggggtgc gctccgcaag 240
ctaagentca tccacttggn nggtttccat tctcggattn cangnagatc aaggnacacc 300
ngntcacanc c 311

<210> 1313

<211> 121
 <212> nucleic acid
 <213> Zea mays

<400> 1313

gctgcccaga aggttggtga ggttgatgct cttgctaagg cattggctgg tcaaaaggat 60
 gaggcttact tcgcagcaaa tgctgctgct caggcctcga ggaaatcctc accccgtgtg 120
 a 121

<210> 1314
 <211> 222
 <212> nucleic acid
 <213> Zea mays

<400> 1314

caccagttca ccacccacct cccaacccca gttcaccccg tcgtcctcgg cgccaccact 60
 cctcgtcccc cggcgtact cccccgtnc acggtccaag gaaagatggc gtcccatatt 120
 gttggatacc ctgcgatggg cccaagagg gagctcaagt ttgccttgga gtctttcttg 180
 gaggggaaga gcagtgccga ggattnggag aaattgcnac tg 222

<210> 1315
 <211> 190
 <212> nucleic acid
 <213> Zea mays

<400> 1315

ccttgattct gacaaattgg ctgcatnntc tgntgcatac gcagnacttg aatctgtact 60
 tntctggattg aatgtgcncg ttgagactta cttngctgnt gttcctgctg agtcnanaa 120
 gaccctaaca ncnngagca gtgtgactgc ntatggtttt gatcttgtcc gtggaacca 180
 nactctnggg 190

<210> 1316
 <211> 120
 <212> nucleic acid
 <213> Zea mays

<400> 1316

gtccacatt gttggatacc ctgcgatggg cccaagagg gagctcaagt ttgcacttga 60

gtctttctgg gatgggaaga acaccgctga agatctggag aaggttgcta ccgacctcag 120

<210> 1317
<211> 159
<212> nucleic acid
<213> Zea mays

<400> 1317

gcttacttcg cagcanatgc tncgtctcat nctctganga nctctcacc ccgtntgacc 60

aatgnagaag tccaaaatgc tgengntgen ctcaanggct atgancancg ccgngggacc 120

aacgtagtg ctagattgga tgnccagcag aagaagctt 159

<210> 1318
<211> 288
<212> nucleic acid
<213> Zea mays

<400> 1318

cgacaccaac atcctctggg tgaaccctga ctgtggtctc aagacacgca agtacacgga 60

ggccaagccc gccctgacca acatggtctc ggccaccaag ctcatccgca ccagcttgc 120

cagcgcgaaa tgaggtcggt tgatagctcc atggtctgat agcgcggaat gagccagttg 180

ttttgaataa tttgggtggt accccctggt ccatggtggt agtgtaggt tagcctotca 240

ttggtgggat acgccgttac aagatgtggt ctaagtttgg agtgtgtg 288

<210> 1319
<211> 268
<212> nucleic acid
<213> Zea mays

<400> 1319

cnagcctnna nagnaacgnt atggttgagt actttcgggt agcagctctc cggttttgcg 60

ttccactgcc aatggctggg tgcagtcnnt atgggtcaag gtgcgtcaag cctccgatca 120

tgctacgggt acgtnagccg ccnaacccc atgaccgttt tctcggtcga agacggcaca 180

gagcatgacc gtctcgccca atgaagggga tgctgactgg ccagtcaca atcctncaac 240

tggtcccttn cgtncgaaa tganccac 268

<210> 1320

<211> 118
 <212> nucleic acid
 <213> Zea mays

<400> 1320

acgacatcca ctgcgccagg atccccctcca ccgaggagat cgcggaccgc gtcaacaaga 60
 tgctcgcngt gctcgacacc aacatcctct ggggtgaaccc tgactgcggt ctcaagac 118

<210> 1321
 <211> 151
 <212> nucleic acid
 <213> Zea mays

<400> 1321

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 ccaggttctt gataccacgg ccattgcttg cgtgttccca gagcgctact cttggactgg 120
 aggtgagatt ggctgagcac tacttctcta t 151

<210> 1322
 <211> 436
 <212> nucleic acid
 <213> Zea mays

<400> 1322

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 tgctcgcngt gctcgacacc aacatcctct ggggtgaaccc tgactgtggt ctcaagacac 120
 gcaagtcacg gaggtcaagc ccgccctgac caacatggtc tnggccacca agcttattcc 180
 gcacccaact tgnacagcgc gaaatgaagt cgtttgatag ctcatggctc gatagcgccg 240
 gaatgagcca agtggtttgn ataatttggg tgttaccccc tgntccatgg ngttagtggg 300
 aaggtagcct ntnattggtg ggataccccc gttccagant nggtctaagg ttggantggg 360
 tggttntctt tnggctatgg ttcttggggg attgtgtgtt ccttgggtat taacnggaat 420
 tgaataccca gtcttt 436

<210> 1323
 <211> 118
 <212> nucleic acid
 <213> Zea mays

<400> 1323

tnatcaccat tgagnnctcn cgggtccgacn agaagctgct ctccgtnttc cgcnaggncg 60
tcaagtagcn cncgggcacn ngcacccggtt tctacnacat ccaactcccc angatccc 118

<210> 1324

<211> 324

<212> nucleic acid

<213> Zea mays

<400> 1324

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ctccccagg atcccgctcg ctgaggagtc gccgaccgca tcgacaagat gtggcgtgct 120
ggacacaaca tctctgggtg aaccccgact gcggctcaag accgcaatac acgagtcaag 180
cccgctgac caacatggtc tccgcccgtc agtcatccgc accagtcgcc agcgccaagt 240
gagcactttt tttgcttttc gttcgaggag ggtgtgatgc aaattgttcn ataaaanggt 300
gtgccgccgt tctgttgtat cggt 324

<210> 1325

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1325

ctctgggtga aacccgactg cggcctcaag acccgcaagt acacggaggt caagcccgcc 60
ctgaccaaca tgggtctccgc cgctaagctc atccgcaccc agctcgccag cgccaagtga 120
gcaccttttt tttgcctttt tcgtttccga ggagggcgtc gtcgatgcca atttgtttcc 180
aataaacagg gctgtgccgc cgttctgttg tactccgtct gtggttaggt tagtagtttt 240
cttgatctcg cccccacgcg gtacctgttt tactactctc tgtttg 287

<210> 1326

<211> 131

<212> nucleic acid

<213> Zea mays

<400> 1326

gtatgccnag cagaagaagc ttaaccnccc catccttccc accanngaca atcggttcgt 60

tcccanaaan cgttgagctc caggaggggc cgccgtgagt ncaaggcaaa aaagatctct 120
gaggaggaat a 131

<210> 1327
<211> 155
<212> nucleic acid
<213> Zea mays

<400> 1327

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tacttcggtg agcagctcct ccggttttgc gttcactgcc aatggctggg tgnagtctta 120
tgggtcaang tgcgtcaagc ctccgatcat ctacg 155

1328
264
nucleic acid
Zea mays
1328

<210> 1328
<211> 264
<212> nucleic acid
<213> Zea mays

<400> 1328
gacgcgtggg tccacaccta tcctgcgcgc tgcgcgtct cctccgtcc cgtcgaccgg 60
aaagatgncg tccacattg ttggataccc tcgcatgggc cccaagaggg agctcaagtt 120
tgcacttgag tctttctggg atgggaagag caccgtgagg attggagaag gttgtacgac 180
ctcaggcagt attggaacag atgntgatnt ggataagtac atcccacaac actttgtnta 240
tacagtcttg cacatgcant gggt 264

<210> 1329
<211> 270
<212> nucleic acid
<213> Zea mays

<400> 1329

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tggtgctctn caaagccgc caaggggtg gagaaggat tccctcttcg ttcctctctt 120
agcagcatcc gccagtgt anaaggaggt cattgctgag ttgaaggcag ctggnggctt 180
catggattca ggttgatgag cccactctng tcctcgnacc tggatgctga caaattgggc 240
tgctttntct gctgcatacg agaantggaa 270

<210> 1330
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 1330

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 ngaggtcant gctgaggtga aggcagctgg ggcttcatgg attcagtttg atganccac 120
 tcttgtcctc gaccttgatt ctgacaaatt ggectgcatt ctctgctgca tacgcagaac 180
 ttgaatctgt actttctgga ttgantgtgc tgtngagann nactttgcng gtgttcctgc 240
 tgantctaca agaccntnta tccntggg 268

<210> 1331
 <211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 1331

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 caacatggtc tccgcgcta agctcatccg cacncagctc gccagcgcca agtgagcacc 120
 ttttttttgc ctttttcgtt tccgaggagg ggtcgtcgat gccaatgtgt ttccaataaa 180
 cagggtgtg nccgcggttc tgttgtactc cgtctgtggt taggttagta nttttcatga 240
 tctcgcnca cgcggtacct gntttactat ctctgtttgg nggttnctga g 291

<210> 1332
 <211> 128
 <212> nucleic acid
 <213> Zea mays

 <400> 1332

 ctncctcctc acctatcctc gccgctgcgc cgtctcctcc gctcccgtcg nccggaaaga 60
 tggccgtccc acattgttgg ataccctcgc atgggccccca agaggagct caagtttgca 120
 cttgagtc 128

<210> 1333
 <211> 286
 <212> nucleic acid

<213> Zea mays

<400> 1333

cacctatcat cgccgctnnc cegtctcttc cgtcccgtc gtnccgaaan atggngtccc 60
acattgntgg atancctcgc atgggccccca aganggagct caagtttgca cttgagtctt 120
tctgggatgg gnagagnacc gctgnaggat ctgganaagg ttgctaccga cntcaagggc 180
cagtancnng angccngatg ggntgatgct gggatcaagt acatccncca gcaacaacct 240
tgctctactt atgacaaggc ccttgacaca ctgccatgct cgcggt 286

<210> 1334

<211> 156

<212> nucleic acid

<213> Zea mays

<400> 1334

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atgctcgccg tgctncgaca ccaacatcct ctgggtgaac cctgactgtg gtctcaagac 120
acgcaagtac acggagggtca agcccgcctt gaacaa 156

<210> 1335

<211> 185

<212> nucleic acid

<213> Zea mays

<400> 1335

ttnangttga tgctcttgct aaggcattgg ctggtcaaaa ggntgaggct tattencagc 60
aaatgctgag ctgaggcttc gaggaatcct caccctgtgt gaaccaatgn agaagtccag 120
aaggctgctg ctgctctcaa gggctctgac caccgctgtg naaccaacgt tatgctagat 180
tggat 185

<210> 1336

<211> 139

<212> nucleic acid

<213> Zea mays

<400> 1336

ggaggttgag gatcttgagg ctgctggtat tcaggatgac cagnttgatg aggcagctct 60

aaggganngt ngccactgcg caagtcccag agcatgcatt ctacctggac tgnngctgtcc 120
actcttttcag antcaccaa 139

<210> 1337
<211> 114
<212> nucleic acid
<213> Zea mays

<400> 1337

cgaggatctt gaggtcgtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 60
agggctgccg ctccgcaagg ctgaacatgc atctaactgg actgggctgt ncca 114

<210> 1338
<211> 254
<212> nucleic acid
<213> Zea mays

<400> 1338

acggagggtca agcccgcctt gaccaacatg gtctccgctg ctaagctcat tcgcacccag 60
ctgccagcgc ccaagtgagc accttttttt gccttttttcg tttccgagga gggcgctcgtc 120
gatgccaaatt tgttttccaat aaacaggggt cccccctgt gcccgccgtt ctgtttgtgt 180
cgtctgtgg ttaggttact agttttcttg atctcgcccc caccggttac ctgttttact 240
atctgttttg tggt 254

<210> 1339
<211> 289
<212> nucleic acid
<213> Zea mays

<400> 1339

cggagggtcaa gcccgccctg accaacaatg tctccgccgc taagctcatc cgcacccagc 60
tcgccagcgc caagtgagca cctttttttt gccttttttcg tttccgagga gggcgctcgtc 120
gatgccaaatt tgttttccaat aaacaggggt gtgccgccgt tctgtttgtac tccgtctgtg 180
gttaggttag tagttttctt gatctcgccc ccacgcggtta cctgtttttac tactctctgt 240
ttggtgggtt ctgaggaagt tgcccgtgta ttgtatcgta aaaaccgag 289

<210> 1340

<211> 252
 <212> nucleic acid
 <213> Zea mays

 <400> 1340

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 ggtctccgcn gctaagctca tcngeacnea gctcgccagc gccaaagtng cacctttttt 120
 ttgccttttt cgtttccgan gagggentcg tngatgccaa tttgtttcna ntaancaggg 180
 ntgtncagc cgttctgttg nactcgtct gtggttagg tagtagtttt cttgatctcg 240
 cccccaagcg nt 252

<210> 1341
 <211> 79
 <212> nucleic acid
 <213> Zea mays

 <400> 1341

 ctgcggagtc caggacacca ccagatcca naccacatg tgtacncaa cttcaacgac 60
 ntctcnant ccatnanng 79

<210> 1342
 <211> 260
 <212> nucleic acid
 <213> Zea mays

 <400> 1342

 cggagggtcaa gcccgccctg accaacaatgg tctccgctgn taagctcatt cgcaccagc 60
 tcgncagcgc caagtgaagca cttttttttg cttttttcgt ttccgaggag ggcgctcgcg 120
 atgccaattt gtttccaata aacagggctc cccccctgtg cccgcngttc tgttgtgtc 180
 cgtctgtggt taggttacta gttttcttga tctcgcccc acgcggtacc tgttttacta 240
 tctgtttggt gggtttctgag 260

<210> 1343
 <211> 124
 <212> nucleic acid
 <213> Zea mays

 <400> 1343

caagggtctc gcaccaccgc cgtgogacca acgttagtgc tagattggat gccagcaga 60
 agangttaac ctccccatcc ttcccaccan tacaatcggg ttcgttccca caaacgtn 120
 agct 124

<210> 1344
 <211> 505
 <212> nucleic acid
 <213> Zea mays
 <400> 1344

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 gaccacgcgc tccggtgaa ctagagtcgg cattctctgg attgaatgtg cttattgaga 120
 catactntgc ngacattcct gctgagtcct acaagaccct cacgtcattg agtgggtgtga 180
 ctgcttacgg ttttgatott atcctgtggag ccaagaccct tgatcttata aggagcagct 240
 tcccctctgg gaagtacctc ttcgctgggtg ttgtggatgg acgcaacatt tgggctgatg 300
 atcttgcgc atctcttagc actcttcagt ctcttgaggc cgttgctggc aaggacaaac 360
 ttgtgggtgc aacctcctgc tcaactgatgc acaccgccgt gatcttgtaa atgagactaa 420
 actggatgaa tgaagattaa gtcattggctt gcatttgctg cccaaaagggt tgcgcaggtt 480
 aatgcccttg cgaagctttg gcaag 505

<210> 1345
 <211> 300
 <212> nucleic acid
 <213> Zea mays
 <400> 1345

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 actaagctgg atgatgagat taagtcatgg cttgcatttg ctgccccaaa ggttggtgag 120
 gttaatgcc ttgccaaggc nttggcaggc caaaaggatg aggtctactt tgcagccaat 180
 gctgctgctc aggcctcaag gagatcatcg cccagggtga caaacganga ggtccagaag 240
 gctgcagctg ctttganggg atctgaccac cgccgttcta ccaactgtntc tgctagattg 300

<210> 1346
 <211> 290
 <212> nucleic acid

<213> Zea mays

<400> 1346

ggacgcaaca tttgggctga tgatcttgct gcatctctta gcaactcttca ttctcttgag 60
gctgttgctg gcaaggacaa acttggtggtg tcaacctcct gctcactgat gcacaccgct 120
gttgaccttg taaatgagac taagctggat gatgagatta agtcatggct tgcatttgct 180
gccccaaaagg ttgttgaggt taatgccctt gccaaaggctt tggcaggcca aaaggatgag 240
gtctactttg cagccaatgc tgctgctcag gctcaagga gatcatcgcc 290

<210> 1347

<211> 259

<212> nucleic acid

<213> Zea mays

<400> 1347

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tgagattaag tcatggcttg catttgctgc ccaaaagggtt gttgagggtta atgcccttgc 120
caaggctttg gcaggccaaa aggatgaggt ctactttgca gccaatgctg ctgtcaggc 180
ctcaaggaga tcatcgccca gggtgacaaa cgaggaggtc cagaaggctg cagctgcttt 240
gaggggatct gaccaccgc 259

<210> 1348

<211> 296

<212> nucleic acid

<213> Zea mays

<400> 1348

tgcttatcga gacatacttc gctgatattc ctgctgagtc ctacaagacc ctcatcatcat 60
tgagtgggtg gactgcttac ggtttcgata ttatccgtgg agccaagacc cttgatctta 120
tcaggagcag cttccctctt gggaagtacc tcttcgctgg tgttgtagat ggacgcaaca 180
tttgggctga tgatcttgct gcatctctta gcaactcttca ttctcttgan gctgttgctg 240
gcaaggacaa acttggtggtg tcaacctcct gctcactgat gcacaccgct gttgac 296

<210> 1349

<211> 272

<212> nucleic acid

<213> Zea mays

<400> 1349

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cttatccgtg gagccaagac ccttgatctt atcaggagca gcttcccctc tgggaagtac 120
ctcttcgctg gtgttggtgga tggacgcaac atttgggctg atgatcttgc tgcattctct 180
agcactcttc agtctcttga ggccgttgc ggcaaggaca aacttggtgt gtcaacctcc 240
tgctcactga tgcacaccgc tgttgatctt gt 272

<210> 1350

<211> 265

<212> nucleic acid

<213> Zea mays

<400> 1350

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gatcttatca ggagcagctt cccctctggg aagtacctct tcgctgggtgt tgtggatgga 120
cgcaacattt gggctgatga tcttgctgca tctcttagca ctcttcagtc tcttgaggcc 180
gttgctggca aggacaaaact tgtggtgtca acctcctgct cactgatgca caccgctggt 240
gatcttgctca atgaaactaa gctgg 265

<210> 1351

<211> 238

<212> nucleic acid

<213> Zea mays

<400> 1351

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cttgaggctg ttgctggcaa ggacaaaact gtggtgtcaa cctcctgctc actgatgcac 120
accgctgttg accttgtaaa tgagactaag ctggatgatg agattaagtc atggcttgca 180
tttgctgccc aaaaggttgt tgaggttaat gcccttgcca aggctttggc aggccaaa 238

<210> 1352

<211> 226

<212> nucleic acid

<213> Zea mays

<400> 1352

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aagctggatg atgagattaa gtcattggctt gcatttgctg cccaaaaggt tgtcgagggt 180
aatgcccttg cgaaggcttt ggcaggccaa aaggatgagg cttact 226

<210> 1353

<211> 248

<212> nucleic acid

<213> Zea mays

<400> 1353

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cttatccgtg gagccaagac ccttgatctt atcaggagca gcttccctc tgggaagtac 120
ctcttcgctg gtgttggtga tggacgcaac atttgggctg atgatcttgc tgcactctt 180
agcactcttc agtctcttga ggcggttgct ggcaaggaca aacttggtgt gtcaacctcc 240
tgctcatg 248

<210> 1354

<211> 203

<212> nucleic acid

<213> Zea mays

<400> 1354

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tcgtctggan cngagtacaa gttcaaggtc tctcncangc ggacttcggc cgctcnaga 120
ttgagctncc cnagttcgaa atgcccgnc ccatgnetnc nccncngatt ncgcnccttn 180
aaancctnnn naggcga aaa ttc 203

<210> 1355

<211> 96

<212> nucleic acid

<213> Zea mays

<400> 1355

ttgggcttna aaactnacct ngnttaagc gcattaccat taaaccccaa acnnancgt 60

tggtgtttcc ccagacnaaa actgngatan anntgc

96

<210> 1356
<211> 518
<212> nucleic acid
<213> Zea mays

<400> 1356

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cccacgcgtc cgcnnacgcg tncgctnacn cgteccgcaca ngcgtcngcn canncgttcg 120

cccacnacgg ncgatnnttc natttccna tctcgtcana ttcaattcgc cgagttcttc 180

ctctctctgc tcgnatggc gctctatgtg ganaanacct cgtcatggac gggagtacaa 240

ggtcaaggat ctatagcang cngacttagn tcgccttgan attgatctgg ccnaggncga 300

aatgcncgna ctcatnggnt gccncgcca gtttgcccg nnaagccct ttggcgntgc 360

taggatctng gnntctcttt acatgaccat tnacaengn gaagtgattg agacnnttac 420

cgngctcgtt gccgaggtnc ngtggtgatc ntcaacatc ttattnacac angaccacgc 480

ttnaancggn attggtncgc ganttggtg canaaatt 518

<210> 1357
<211> 447
<212> nucleic acid
<213> Zea mays

<400> 1357

cggcaagatc cccgaccgg agtccaccga caacgctgag ttcaagatcg tgctcacnat 60

catccgcgac gggctcaagg ctgaccccaa gaagtaccgc aagatgaagg agaggcttgt 120

cggcgtctct gaggagacca ccacgggtgt caagaggctc taccagatgc aggagaccgg 180

cgcctctctc ttccctgcca ttaacgtcaa cgattccgtc accaagagca agtttgacaa 240

cctgtatggt tgccgccact cgtccctga tggctgatg agggccactg acgttatgat 300

cgccgaaaag gttgccgtgg tctgcggata cggatgatgc ggcaagggtt gtgctgctgc 360

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gctctgatgg anggtcttta ngtcctt 447

<210> 1358

<211> 515
 <212> nucleic acid
 <213> Zea mays

<400> 1358

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gnnnagnnnn nnnannnttt gnangggggg gggaagagan attttancgg tccggattcc 60
cgggtcgacc cacgcgtccg ccatcatccg cgacgggctc aaggctgacc ccaagaagta 120
ccgcaagatg aaggagaggc ttgtcggcgt ctctgaggag accaccacgg gtgtcaagag 180
gctctaccag atgcaggaga cgggcgccct cctcttcctt gccattaacg tcaacgattc 240
cgtcaccaag agcaagtttg acaacctgta tggttgccgc cactcgctcc ctgatgggtc 300
gatgagggcc actgacgtta tgatcgccgg taaggttgct gtggtctgcg gatacgggtga 360
tgtcggcaag ggttgtgtg ctgccctcaa gcaggctggt gcccggtgtca ttgtgaccga 420
gatcgacccc atctgtgccc tccaggctct gatggaaggg tcttcagggtc cttcccttgg 480
aagacgttgt ctctgaactg acatcttctg gacca 515

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<210> 1359
 <211> 530
 <212> nucleic acid
 <213> Zea mays

<400> 1359

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ggnnnagggt gntttngtcc naanntntnt tttncttctg ccngtaccgg tccggattcc 60
cgggtcgacc cacgcgtccg cccacgcgtc cgtggttgac cacatgagga agatgaagaa 120
caatgccatt gtctgcaaca ttggccactt tgacaatgaa attgatatgc tcggccttga 180
gacctaccct ggcgtcaagc gcataccat caagccccag actgaccgct ggggtgtccc 240
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tactggccat cctagctttg tcatgtctg ctcatcact aaccaggtea ttgcccact 360
tgaactgtgg aaggagaaga gctctggcaa gtatgagaag aagggtgatg tgcctcccaa 420
gcaccttgat gagaagggtg ctgctctcca cttgggcaag cttggtgcca agctgaccaa 480
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<210> 1360
 <211> 501
 <212> nucleic acid

<213> Zea mays

<400> 1360

ggnngnnnngn aattctaacn tncngcagcc tgcanaagtc canggtccac ccacncgtcg 60
ncgaggagga gtacgagaag accggcaaga tccccgaccc ggagtccacc gacaacgcga 120
gttcaagatc gtgtcacca tcatccgga cgggctcaag gctgaccca agaagtacgc 180
aagatgaagg agaggcttgt cggcgtctct gaggagacca ccacgggtgt caagagctct 240
accagatgca ggagaccggc gccctcctct tccctgccat taacgtcaac gattcgtcac 300
caagagcaag tttgacaacc tgtatgggtg ccgccactca ctccctgatg gtctatgagg 360
gccaccgacg ttatgatcgc cggtaagggt gccgtggtct gcggatacgg tgagttggca 420
agggttgtgc cgctgcactc aaancangct gggtgcccgt gtcaantgtg acgaagattt 480
aaccaatct gcgncctcca a 501

<210> 1361

<211> 473

<212> nucleic acid

<213> Zea mays

<400> 1361

cgccgnaag gttgctgtgg tctgcggata cggatgatgc ggcaaggggt gtgctgctgc 60
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ngctctgatg gagggctctc angtccttcc cttggaggac gttgtctctg aagctgacat 180
cttcgtgacc accactggca acaaggatat catnatggat gaccacatga ggaagatgaa 240
gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcgggct 300
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nccccgagac caacactggg catcattgtc cttgctgaag gtcggctgat naaccttgng 420
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<210> 1362

<211> 524

<212> nucleic acid

<213> Zea mays

<400> 1362

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 gccatggcgc tctcngtgga gaagacctcg tctggacggg agtacanggt caaggatctc 180
 tcgcaggcgg acttcggccg cctcgagatt gagctggccg aggtcgaaat gcccggcctc 240
 atggcgtgcc gcgccgagtt cggcccgctc aagcccttcg ccggcgctag gatctcgggg 300
 tctctccaca tgaccatcca ngaccggcgt cctcatcgag accctcaccg cgctcggcgc 360
 cgangtccgc tgggtctcct gcaacatctt ctccacgcag gaccacgccg ccgccgccat 420
 cgcgcgcgac tcggccgccg tgttcgcctg gaanggggaa accctcgang agtactggtg 480
 gtgcancgaa cgctgcctcg actggggggc angcgggccg cccc 524

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<210> 1363
 <211> 478
 <212> nucleic acid
 <213> Zea mays
 <400> 1363
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 tccatttccc catctcccag atccaattcg cgagttctcc ctctcttgcc gccatggcct 120
 ctctgtggag aagacctcgt ctggacggga gtacaaggtc aaggatctct cgcaggcgac 180
 ttcggccgcc tcgagattga gctggccgag gtcgaaatgc ccggcctcat ggctgtcgcg 240
 ccgagttcgg cccgtcgaag cccttcgctg gcgctaggat ctcggggtct ctccaatgac 300
 catccagacc gccgtctcct tcgagaccct caccgcgctc ggccgccgang tccgtggtgc 360
 tcctgcaaca tttctccac gcaggaccac gccgccgccg ccctcgcgcg cgatcggccg 420
 ccgtgttcgc ctggaagggg gaagacctcg aggagtactg gtggtgcacc gacgctgc 478

<210> 1364
 <211> 528
 <212> nucleic acid
 <213> Zea mays
 <400> 1364
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 anttcccggg tcgaccacg cgtccgcccc atctcccaga tccaattcgc gagttctttn 120

tcctctgccg ccatggcgct ctctgtggag aagacctcgt ctggacggga gtacaaggtc 180
aaggatctct cgcaggcgga cttcggccgc ctcgagattg agctggccga ggtcgaaatg 240
cccggcctca tggcgtgccg cgcgcagttc ggcccgtcca agcccttcgc cggcgctagg 300
atctcggggt ctctccacat gaccatccag accgcgctcc tcctcgagac cctcaccgcg 360
ctcggcgccg aggtccgctg gtgctcctgc aacatcttct ccacgcagga ccacgccgcc 420
gccgccatcg cgcgcgactc ggccgccgtg ttgccttga agggggagac cctcgaggag 480
tactggtggt gcacccgagc gctgnctnga ctggggcnag gcggggccg 528

<210> 1365
<211> 354
<212> nucleic acid
<213> Zea mays

<400> 1365
cgtgcgcgcg cgagttcggc ccgtccaagc ccttcgccgg cgctaggatc tcggggtctc 60
tccacatgac catccagacc gccgtcctca tcgagaccct caccgcgctc ggcgcgcagg 120
tccgctggtg ctcttgcaac atcttctcca cgcaggacca cgccgccgcc gccatcgcg 180
gcgactcggc cgcctgttcc gcctggaagg gggagaccct cgaggagtac tgggtggtgca 240
ccgagcgctg cctcgactgg ggcgaggcgg gcggccccga cctcatcgtc gacgacggcg 300
gcgacgccac gctgctcatc cacgagggtg tcaaggccga ggaggattac nana 354

<210> 1366
<211> 476
<212> nucleic acid
<213> Zea mays

<400> 1366
gggggngnaa cttcanntcc cagnngccgg tcaagaatta acgggtcgac gcncgcgtcc 60
ggctcaaggc tgaccccaag aagtaccgca agatgaagga gaggcttgct ggctctctct 120
aggagaccac cacgggtgtc aagaggctct accagatgca ggagaccggc gccctcctct 180
tccctgccat taacgtcaac gattccgtca ccaagagcaa gtttgacaac ctgtatggtt 240
gccgccactc gtcctctgat ggtctgatga gggccactga cgttatgatc gccggaaagg 300
ttgccgtggt ctgcggatac ggtgatgtcg gcaagggttg tgctgctgcc ctcaagcagg 360

ctggtgcccg tgtcattgtg accgagatcg accccatctg tgcctccag gctctgaatg 420
 ganggtcttc aagtccttcc cttggaggac gttgtctcct gaagctgnca tcttng 476

<210> 1367
 <211> 468
 <212> nucleic acid
 <213> Zea mays

<400> 1367

atcatggttg accacatgag gaagatgaag aacaatgcc a ttgtctgcaa cattggccac 60
 tttgacaatg aaattgatat gctcggcctt gagacctacc ctggcgtcaa gcgcatcacc 120
 atcaagcccc agactgaccg ctgggtgttc cccgagacca aactggcat cattgtcctt 180
 gctgagggtc gcctgatgaa ccttgggtgt gctactggcc atcctagctt tgtcatgtcc 240
 tgctcattca ctaaccaggt cattgcccc a cttgaactgt ggaaggagaa gagctctggc 300
 aagtatgaga agaaggtgta tgtgcttccc aagcaccttg atgagaaagg tgctgtctc 360
 acttggggca agctttggtg ccaagcttac caaacttaac caagtcttaa ggccgactac 420
 attaaccgtg ccgcatcgaa gggtccttac aagcctgcc actnaccg 468

<210> 1368
 <211> 484
 <212> nucleic acid
 <213> Zea mays

<400> 1368

tttctttgcc ctggccggta ccanatcana gaattcccgg gncgaccac gcatccngct 60
 caaggctgac cccaagaagt accgcaagat gaaggagagg cttgtcggcg totctgagga 120
 gaccaccacg ggtgtcaaga ggctctacca gatgcaggag accggcgccc tcctcttccc 180
 tgccattaac gtcaacgatt ccgtcaccaa gagcaagttt gacaacctgt atggttgccg 240
 cactcgctc cctgatggtc tgatgagggc cactgacgtt atgatcgccg gaaaggttgc 300
 cgtggctctgc ggatacgggtg atgtcggcaa gggttgtgct gctgccctca agcaggctgg 360
 tgcccgtgtc attgtgaccg agatcgaccc catctgtgcc tccanggctc tgaatggaag 420
 gtcttcangt ccttcccttg gaaggacgtt ggtctccgaa agctgacatc ttcgtgacna 480
 acaa 484

<210> 1369
 <211> 416
 <212> nucleic acid
 <213> Zea mays

<400> 1369

tgctcggcct tgagacctac cctggcggtca agcgcatcac catcaagccc cagactgacc 60
 gctgggtggtt ccccgagacc aacactggca tcattgtcct tgctgaggggt cgcctgatga 120
 accttgggtg tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg 180
 tcattgcccc acttgaactg tggaaggaga agagctctgg caagtatgag aagaagggtgt 240
 atgtgtcccc caagcacctt gatgagaagg ttgctgtctt ccacttgggc aagcttggtg 300
 ccaagctgac caagctcacc aagtctcagg ccgactacat cagcgtgccg atcgaggggc 360
 cctacaagcc tgccactacc ggtactaggc aagccaggca cacggcttgc agctna 416

<210> 1370
 <211> 452
 <212> nucleic acid
 <213> Zea mays

<400> 1370

attgatatgc tcggccttga gacctaccct ggcgtcaagc gcatcaccat caagccccag 60
 actgaccgct ggggtgttccc cgagaccaac actggcatca ttgtccttgc tgaggggtgc 120
 ctgatgaacc ttgggtgtgc tactggccat cctagctttg tcatgtcctg ctcattcact 180
 aaccagggtca ttgcccact tgaactgtgg aaggagaaga gctctggcaa gtatgagaag 240
 aagggtgtatg tgctcccca gacacttgat gagaagggtg ctgctctcca cttggggcaa 300
 gcttggtgcc aagctgacca agctcaccaa gtcttaagcc gactacatca gcgtgccgat 360
 cganggtccc tacaagcctg ccactaccgg tactangcaa gccagcacac ggcttgcagc 420
 ttactcggcc cgttgtgtgc tatgaagttc ct 452

<210> 1371
 <211> 481
 <212> nucleic acid
 <213> Zea mays

<400> 1371

gcaagatgaa ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggn 60
nctaccagat gcaggagacc ggcgccctcc tnttcctgc cattaacgtc aacgattccg 120
tcaccaagag caagtttgac aacctgtatg gntgccgcca ctgcgtccct gatggtctga 180
tgagggccac tgacgttatg atcgccggtg aggntgctgn ggtctgcgga tacgngatg 240
tcggcaaggg ttgngctgct gccctcaagc aggctggtgc cegtgtcatt gggaccgaga 300
tcgaccccat ctgngccctc caaggctctg atggagggnc ttcaggncct tcccttgag 360
gacgnngctn tgaaactgac atcttcggga ccaccactgg caacaaggat atcatcangg 420
gtgaccacat gaaggaagat gaagaacaaa gccaanngcn gggaacaatn ggccaanttg 480
a 481

<210> 1372
<211> 445
<212> nucleic acid
<213> Zea mays

<400> 1372
cgccgtctc atcgagacc tcaccgcgt cggcgccgag gtcgctggt gtcctgcaa 60
catcttctcc acgcaggacc acgccgccgc cgccatcgcg cgcgactcgg ccgcgtggt 120
cgctggaag ggggagacc ttgaggagta ctggtggtgc accgagcgt gctcgactg 180
ggcgaggcg ggcggcccg acctcatcgt cgacgacggc ggcgaccca cgctgctcat 240
ccacgagggt gtcaaggccg aggaggacta cgagaagacc ggcaagatcc ccgatccgga 300
gtcaccgaca acgctgagtt caagatcgt ctnaccatca ttccgcgaan ggctnaaagn 360
ttaccccaag aagtcccga aaggataaag aaaagcttgt cggcgtnttt taagagacca 420
cacnggtgtn aaaaggnttt acaga 445

<210> 1373
<211> 512
<212> nucleic acid
<213> Zea mays

<400> 1373
ggggttggtg nnnnggtggg naaattatnt gttgtaccgg tccggaattc ccgggtcgac 60
ccacgcgtcc gccacgcgt ccgctcacgc gttcgccan nntccagat ncaattcgcg 120

agttctgggn tgctctgccg ccatggcgct ctctgtggag aagacctcgt ctggacggga 180
gtacaaggtc aaggatctct cgcaggcgga cttcggtcgc ctcgagattg agctggccga 240
ggtcgaaatg cccggcctca tggcgctgcc cgccgagttc ggcccgtcca agcccttcgc 300
cggcgctagg atctcggggg ctcttcacat gaccatccag accgccgtcc tcatcgagac 360
cctcaccgcg ctcggcgcgc aggtccgctg gtgctcctgc aacatcttct ncacgcagga 420
ccacgccgtc ggcggcacgc ngcgcgactc gngcggcggtg ttgccttgga agggggagac 480
cctttganga gtactggtgg tncaccnaac cg 512

<210> 1374
<211> 523
<212> nucleic acid
<213> Zea mays

<400> 1374
agaggctngt tttgantggg gattaanggn ncttaatgg aaagctcttc cggnccggn 60
nccccgctcg acccangcnn tncanntnnt ggaggggtctt naggtcctnc ncttggagga 120
cgttgtctnn gnagctgaen gnttcgtgac caccactggc aacaaggata tcatcatggt 180
tgaccacatg aggaagatga agaacaatgc cattgtctgc aacattggcc actttgacaa 240
tgaaattgat atgctcggcc ttgagacctc ccttggcgtc aagcgcatna ccatcaagcc 300
ccagactgac cgctgggtgt tccccgagac caacactggc atcattgtcc ttgctgaggg 360
tcgcctgatg aaccttgggt gtgctactgg ccatcctagc tttgtcatgt cctgctcatt 420
cactaaccag gtcattgccc aacttgaact gtggaaggag aagagctctg gcaagtatga 480
gaaagaangt gtatgtgctt cccaacacct ttgatgagaa ggt 523

<210> 1375
<211> 441
<212> nucleic acid
<213> Zea mays

<400> 1375
cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60
cctcgtctgg acnggagtac aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120
agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gaggctcgcc 180

cgccaagcc cttcgccggc gctaggatct cggggtctct ccacatgacc atccagaccg 240
 ccgtctcat cgagaccctc accgcgctcg gcncccgagg tccgctgggtg ctcttgcaac 300
 atcttctnca cgcaggacca cnccttcgnc ngcatcgcg cgcactcgnc cgcctgtgtc 360
 gcctggaaag ggggagnccc tcnaggagta ctgggtgttc accnagccgc ttgccttgnc 420
 tggggccnag gcngnccgnc c 441

<210> 1376
 <211> 488
 <212> nucleic acid
 <213> Zea mays

<400> 1376

ggnnnnnngg naacttnaac tncagcgcc aggtaacggt caaagaattc ccgggtcgac 60
 caccggtccg cggacgcgtg ggcccgcttc atttcccat ctcccagatc caattcgag 120
 ttctccctcc tctgcgcga tggcgctctc tgtggagaag acctcgtctg gacgggatac 180
 aaggtaagg atctctcgca ggcggacttc ggccgcctcg agattgagct ggccgagtcg 240
 aaatgcccg cctcatggcg tgccgcgcg agttcggccc gtccaagccc ttgcgggcgc 300
 taggatctcg gggctctctc acatgaccat ccagaccgcc gtcctcatcg agacctcacc 360
 gcgctcggcg ccgaggtccg ctgggtgctc tgcaacatct tctccacgca aggccacgcc 420
 gccgcgcga tgcgcgcga ctggncgcc gtgttcgcct ggaaggggga gactcgaagg 480
 attatggg 488

<210> 1377
 <211> 325
 <212> nucleic acid
 <213> Zea mays

<400> 1377

caagaagtac cgcaagatga aggagaggct tgtcggcgtc tctgaggaga ccaccacggg 60
 tgtcaagagg ctctaccaga tgcaggagac cggcgccctc ctcttccctg ccattaacgt 120
 caacgattcc gtcaccaaga gcaagtttga caacctgtat ggttgccgcc actcactccc 180
 tgatggtctg atgagggcca ccgacgttat gatcgccggt aaggttgccg tggctctgcg 240
 atacggtgat gttggcaagg gttgtgccgc tgcactcaag caggctgggtg ccggtgtcat 300

tgtgaccgag atcgacccca tctgc 325

<210> 1378
 <211> 325
 <212> nucleic acid
 <213> Zea mays

<400> 1378

gaagctgaca ttttctgtgac caccactggc aacaaggata tcatcatggt tgaccacatg 60
 aggaagatga agaacaatgc cattgtctgc aacattggcc actttgacaa tgaaattgat 120
 atgctogggc ttgagaccta ccctggcgtc aagcgcatca ccatcaagcc ccagactgac 180
 cgttggtgtg tccccgagac caacactggc atcattgtcc ttgctgaggg tcgcctgatg 240
 aacottgggt gtgctactgg ccctcctagc tttgtcatgt cctgtcatt cactaaccag 300
 gtcattgccc aacttgaact gtgga 325

<210> 1379
 <211> 370
 <212> nucleic acid
 <213> Zea mays

<400> 1379

gtggtgcacc gagcgctgcc tcgactgggg cgaggcgggc ggccccgacc tcatcgtcga 60
 cgacggcggc gacgcancgc tgctacatcc acgagggtgt caaggaccga ggaggattac 120
 gagaagaccg gcaagatccc cgacccggag tccaccgaca acgctgagtt caagatcgtg 180
 ctcaccatca tccgcgacgg gctcaaggt gacccaaga agtntcgcaa gatgaaggag 240
 aggettgtcg gcgtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag 300
 gagaccggcg ccctcctctt ccctgccatt aacgtcaaga ttccgtcacc aagagcaatt 360
 tgacaacctg 370

<210> 1380
 <211> 329
 <212> nucleic acid
 <213> Zea mays

<400> 1380

gcaagtttga caacctgtat ggttgccgcc actcgctccc tgatgggtctg atgagggcca 60

ctgacgttat gatcgccgga aaggttgccg tggctctgcg atacggtgat gtcggcaagg 120
gttgtgctgc tgccctcaag caggctgggtg cccgtgtcat tgtgaccgag atcgacccca 180
totgtgeect ccaggctctg atggaggggc ttcaggtcct tcccttgag gacgttgtct 240
ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg gttgaccaca 300
tgangaagat gaagaacaat gccattgtc 329

<210> 1381
<211> 346
<212> nucleic acid
<213> Zea mays

<400> 1381

cacgggtgtc aagaggctct accagatgca ggagaccggc gccctcctct tccctgccat 60
taacgtcaac gattccgtca ccaagagcaa gtttgacaac ctgtatgggt gccgccactc 120
actccctgat ggtctgatga gggccaccga cgttatgatc gccggtaagg ttgccgtgggt 180
ctgoggatac ggtgatgttg gcaaggggtg tgccgctgca ctcaagcagg ctggtgcccg 240
tgtcatgtga ccgagatcga ccccatctgc gccctccagg ctctgatgga ggggtcttcag 300
tccttccctt ggaggacgtt gtctcggaag ctgacatctt cgtgac 346

<210> 1382
<211> 320
<212> nucleic acid
<213> Zea mays

<400> 1382

ccgagcgtg ccttgactgg ggcgaggcgg gcggccccga cctcatcgtc gacgacggcg 60
ggaagccac gctgctcatc cacgagggtg tcaaggccga ggaggagtac gagaagaccg 120
gcaagatccc cgacnccgag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 180
tccgcgacgg gctcaaggct gaccccaaga agtaccgcaa gatgaaggag aggcttgtcg 240
gcggtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 300
ccctcctctt cctgccatt 320

<210> 1383
<211> 455
<212> nucleic acid

<213> Zea mays

<400> 1383

ctcaaggctg accccaagaa gtaccgcaag atgaaggaga ggcttgtcgg cgtctctgag 60
gagaccacca cgggtgtcaa gaggtcttac cagatgcagg agaccggcgc cctcctcttc 120
cctgccatta acgtcaacga ttccgtcacc aagagcaagt ttgacaacct gtatggttgc 180
cggcactcgc tcctgatgg tctgatgagg gccactgacg ttatgatcgc cggtaagggtt 240
gctgtggtct gcggatacng tgatgtcggc aagggttgtg ctgctgcctn aaacaggctg 300
gtgccccgtg tcattgtgac ccagatcgac cccatctgtg cccttcaagc ttctgatnga 360
nngncttcan gtccttcctt tggaaggacg ttgtntttgn aacttgacat ttttngntgg 420
accaccactt gggaacaagg ggtnttatta ttggg 455

<210> 1384

<211> 317

<212> nucleic acid

<213> Zea mays

<400> 1384

cgagatcgac cccatctgtg ccctccaggc tctgatggag ggtcttcagg tccttccctt 60
ggaggacgtt gtctctgaag ctgacatctt cgtgaccacc actggcaaca aggatatcat 120
catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 180
tgacaatgaa attgatatgc tcggccttga gacctaccct ggcgtcaagc gcatcaccat 240
caagccccag actgaccgct ggggtgttccc cgagaccaac actggcatca ttgtccttgc 300
tgagggtcgc ctgatga 317

<210> 1385

<211> 332

<212> nucleic acid

<213> Zea mays

<400> 1385

gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct 60
tgagacctac cctggcgtca agcgcacac catcaagccc cagactgacc gctgggtgtt 120
cccagacca aactggcat cattgtcctt gctgagggtc gcctgatgaa ccttgggtgt 180

gctactggcc atcctagctt tgtcatgtcc tgctcattca ctaaccaggt cattgccc aa 240
 cttgaactgt ggaaggagaa gagctctggc aagtatgaga agaagggtga tgtgctcccc 300
 aagcaccttg atgagaaggt tgctgctctc ca 332

<210> 1386
 <211> 337
 <212> nucleic acid
 <213> Zea mays

<400> 1386

cggcgcctc ctcttccctg ccattaacgt caacgattcc gtcaccaaga gcaagtttga 60
 caacctgtat gggttgcgcc actcactccc tgatgggtctg atgagggcca ccgacgttat 120
 gatcgccggt aaggttgcgc tgggtctgcgc atacgggtgat gttggcaagg gttgtgcgc 180
 tgcaactcaag caggttgggtg cccgtgtcat tgtgaccgag atcgaccca tctgcgcct 240
 ccaggtcttg atggaggggtc ttcaggtcct tcccttgag gagttgtctc ggaagctgac 300
 atcttcgtga ccacccatgg caacaaggat atcatca 337

<210> 1387
 <211> 316
 <212> nucleic acid
 <213> Zea mays

<400> 1387

gtcttcaggt ccttcccttg gaggacgttg tctctgaagc tgacatcttc gtgaccacca 60
 ctggcaacaa ggatatcatc atggttgacc acatgaggaa gatgaagaac aatgccattg 120
 tctgcaacat tggccacttt gacaatgaaa ttgatatgct cggccttgag acctaccctg 180
 gcgtcaagcg catcaccatc aagccccaga ctgaccgctg ggtgttcccc gagaccaaca 240
 ctggcatcat tgtccttgct ganggtcgcc tgatgaacct tgggtgtgct actggccatc 300
 ctagctttgt catgtc 316

<210> 1388
 <211> 315
 <212> nucleic acid
 <213> Zea mays

<400> 1388

gagctcggcg ccgaggtccg ctggtgctcc tgcaacatct tctccacgca ggaccacgcc 60
gcccgcgcca tncgcgcna ctcgcccgcc gtgttcgcct ggaaggggga gacccttgan 120
gagtactggg ggtgcaccga gcgctgcctt gactggngcg angcggggcg ccccnacctc 180
atcgctgcag acggcggcga cgccacgctg ctcatccaag aggggtgtcaa ggccgaggag 240
gagtacgaga agaccggcaa gatccccgac ccggagtcca ccgacaacgc tgagttcaag 300
atcgtgggtca ccatac 315

<210> 1389
<211> 310
<212> nucleic acid
<213> Zea mays

<400> 1389

tctgaggaga ccaccacggg tgtcaagagg ctctaccaga tgcaggagac cggcgcctc 60
ctcttccctg ccattaacgt caacgattcc gtcaccaaga gcaagtttga caacctgtat 120
ggttgcgcgc actcgtctcc tgatggtctg atgagggcca ctgacgttat gatcgccgga 180
aaggttgcgc tgggtctgcg atacggtgat gtcggcaagg gttgtgctgc tgccctcaag 240
caggctggtg cccgtgtcat tgtgaccgag atcgacccca tctgtgccct ccaggctctg 300
atggaggggc 310

<210> 1390
<211> 457
<212> nucleic acid
<213> Zea mays

<400> 1390

ggggtctctc cacatgacca tccagacnng caagatcccc gatccggagt ccaccngna 60
cgctgagttc aagatcgtgc tcaccatcat ccgcgacggg ctcaaggctg accccaagan 120
gtaccgcaag atgaaggaga ggcttgctcg cgtctntgag gagaccacca cgggtgtcaa 180
gaggctctac cagatgcagg agaccggcgc cctcctcttc ctgccattaa cgtcaacgat 240
tccgtcacca agagcaagtt tgacaacctg tatggttgcc gncactcgct ccctgatggg 300
ctgatgaagg gccactgacc ttatgatcgc ccgaaanggt gccgtgggtc gcggataccg 360
tgatgtcngc aaagggttgt gcttnttnan ttaaancang cttggtggcc ctgtcantnt 420

gaaccananc caancccatn ttggncctt cagggtt

457

<210> 1391
<211> 520
<212> nucleic acid
<213> Zea mays

<400> 1391

agttnattna aggnngngg ntgnaagnta tactcgtncg gaattcccggt gtcgacccac 60
gcggtccgac tncagaccc aattcngag ttctnnntan tctgccgaca tggcgctctc 120
tgtggagaag acctgntctg gacgggagta caaggtcaag gatctctagc aggcggactn 180
aggccgacta gagattgagc tggccgaggt cgaaatgcc ggccatnatgg cgtgccgngc 240
cgagttcggc ccgtncgaagc ccttngccgg cgctaggatc tcgggggtctc tccacatgac 300
cattcagacc gncgtcctca tcgagaccct caccgcgctc ggcgccgagg tccgctggtg 360
ctcctgcaac atctttcttc acgcangacc acgccgncgn cgtcatcgcn cgcgactcgg 420
ccggcgtggt cgcttggaaan ggggagaccc ttgangagtc tgggtggtgca ccgnacgctt 480
gcntanntgg gccaaaggcc cggcccgacc tattgtngac 520

<210> 1392
<211> 305
<212> nucleic acid
<213> Zea mays

<400> 1392

cgtcgacgac ggccggcgacg ccacgtgct catccacgag ggtgtcaagg ccgaggagga 60
ttacgagaag accggcaaga tctccgaccc ggagtccacc gacaacgctg agttcaagat 120
cgtgctcacc atcatccggc acgggctcaa ggctgacccc aagaagtacc gcaagatgaa 180
ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 240
gcaggagacc ggcgccctcc tcttccctgc cattaacgct aacgattccg tcaccaagag 300
caagt 305

<210> 1393
<211> 317
<212> nucleic acid
<213> Zea mays

<400> 1393

gtgcccgtgt cattgtgacc gagatcgacc ccatctgcnc cctccaggct ctgatggagg 60
gttttcaggt ctttcccttg gaggacgttg tctcggaagc tgacatcttc gtgaccacca 120
ctggcaacaa ggatatcatc atggttgacc acatgaggaa gatgaagaac aatgccattg 180
totgcaacat tggccacttt ganaatgaaa ttgatatgct cggccttgag acctaccctg 240
gcgtaagcg catcaccatc aagccccaga ctgaccgctg ggtgttcccc gagaccaaca 300
ctggcatcat tgtcctt 317

<210> 1394

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1394

gctgccttga ctggggcgag gcgggcgggc ccgacctcat cgtcgacgac ggcggcgacg 60
ccacgctgct catccacgag ggtgncaagg ccgaggagga gtacgagaag accggcaaga 120
tccccgaacc ggagtccacc gacaacgctg agttcaagat cgtgctcacc atcatccgcg 180
acgggctcaa ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct 240
ctgaggagac caccacgggt gtcaagaggc tctaccagat gcaggagacc ggcgccctcc 300
tcttccctg 309

<210> 1395

<211> 335

<212> nucleic acid

<213> Zea mays

<400> 1395

atcaagcccc agactgaccg ctgggtgttc cccgagacca aactggcat cattgtcctt 60
gctgagggtc gctgatgaa ccttgggtgt gctactggcc atcctagctt tgtcatgtcc 120
tgtcattca ctaaccagggt cattgcccaa cttgaactgt ggaaggagaa gagctctggc 180
aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgtgctctc 240
caattgggca agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 300
agcgtgccga tcgaggggtcc ctacaagcct gccca 335

caacctgtat ggttgccgcc actcgctccc tgatggtctg atgagggcca ctgacgttat 240
 gatcgccgga aaggttgccg tggctctgcg atacggtgat gtcggcaagg gttgtgctgc 300
 tgc 303

<210> 1399
 <211> 311
 <212> nucleic acid
 <213> Zea mays

<400> 1399

agcaggctgg tgcccgtgtc attgtgaccg agatcgaccc catctgcnc ctcaggctc 60
 tgatggaggg tcttcaggct cttcccttgg aggacgttgt ctcggaagct gacatcttcg 120
 tgaccaccac tggcaacaag gatatcatca tggttgacca catgaggaag atgaagaaca 180
 atgccattgt ctgcaacatt ggccactttg acaatgaaat tgatatnctc ggcccttgaga 240
 cctaccctgg cgtcaagcgc atcaccatca agccccagac tgaccgctgg gtgttccccg 300
 agaccaacac t 311

<210> 1400
 <211> 308
 <212> nucleic acid
 <213> Zea mays

<400> 1400

caaggatata atcatggttg accacatgag gaagatgaag aacaatgcca ttgtctgcaa 60
 cattggccac tttgacaatg aaattgatat gtcggcctt gagacctacc ctggcgtcaa 120
 ggcacacacc atcaagcccc agactgaccg ctgggngttc cccgagacca aactggcat 180
 cattgtcctt gctgagggct gctgatgaa cttgggtgt gctactggcc atcctagctt 240
 tgtcatgtcc tgctcattca ctaaccaggt cattgccc aa cttgaactgt ggaaggagaa 300
 gagctctg 308

<210> 1401
 <211> 309
 <212> nucleic acid
 <213> Zea mays

<400> 1401

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 ggagggtott cagggtccttc ccttgaggga cgttgtctcg gaagctgaca tcttcgtgac 120
 caccactggc aacaaggata tcatcatggt tgaccacatg aggaagatga agaacaatgc 180
 cattgtctgc aacattggcc actttgacaa tgaaattgat atgctcggcc ttgagacctt 240
 ccctggcgtc aagcgcatca ccatcaagcc ccagactgac cgctgggtgt tccccgagac 300
 caacactgg 309

<210> 1402
 <211> 311
 <212> nucleic acid
 <213> Zea mays

<400> 1402

ctctgtgacc accactggca acaaggatat catcatgggt gaccacatga ggaagatgaa 60
 gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct 120
 tgagacctac cctggcgta agcgcatcac catcaagccc cagactgacc gctgggtggt 180
 ccccgagacc aacaactggca tcattgtcct tgctgagggt cgcctgatga accttgggtg 240
 tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg tcattgcca 300
 cttgaactgt g 311

<210> 1403
 <211> 338
 <212> nucleic acid
 <213> Zea mays

<400> 1403

gtttgacaac ctgtatgggt gccgccactc gctccctgat ggtctgatga gggccactga 60
 cgttatgatc gccggaaang ttgccgtggt ctgcggatac ggtgatgtcg gcaagggttg 120
 tgctgctgcc ctcaagcagg ctggtgcccg tgtcattgtg accgagatcg acccatctg 180
 tgccctccag gctctgatgg agggctttca ggtccttccc ttggaggacg ttgtctctga 240
 agctgacatc ttctgacca ccaactggcaa caaggatatc atcagggtga ccacatgang 300
 aagatgaaga acaatgccat gtctgcaaca tggccant 338

<210> 1404

<211> 306
 <212> nucleic acid
 <213> Zea mays

 <400> 1404

 ggagacctac cctggcgtca agcgcacac catcaagccc cagactgacc gctgggtgtt 60
 ccccgagacc aacactggca tcattgtcct tgctgagggt cgctgatga accttgggtg 120
 tgctactggc catcctagct ttgtcatgtc ctgtcattc actaaccagg tcattgccc 180
 acttgaactg tggaaggaga agagctctgg caagtatgag aagaagggtg atgtgctccc 240
 caagcacctt gatgagaagg ttgctgctct ccacttgggc aagcttgggtg ccaagctgac 300
 caagct 306

<210> 1405
 <211> 424
 <212> nucleic acid
 <213> Zea mays

 <400> 1405

 anangetncln gcggtctcct gntacaanag annnnnnnaa nggttgngct natgccgtna 60
 ancaanntga ngcccagatc attgtnaccg agatcgaccc catctgtgcc ctncagggtc 120
 tgatggaggg tcttcagggtc cttcccttgn aggacgttgt ctctgaagct gacatcttcg 180
 tgaccaccac tggcaacaag gatatcatca tggttgacca catgaggaag atgaagaaca 240
 atgccattgt ctgcaacatt ggccactttg acaatgaaat tgatatgctc ggccttgaga 300
 cctaccctgn cgtcaagcgc atcaccatca agccccagac tgaccgctgg gtgttccccg 360
 agaccaacac ttggcattca ttgtccttgc tgaaggtcnc cctgattaac ctttggttgt 420
 gcta 424

<210> 1406
 <211> 299
 <212> nucleic acid
 <213> Zea mays

 <400> 1406

 gtgaccgaga tcgaccccat ctgtgccctc caggctctga tggaggggtct tcaggctcctt 60
 cccttggagg acgttgtctc tgaagctgac atcttcgtga ccaccactgg caacaaggat 120

atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 180
cactttgaca atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcac 240
accatcaagc cccagactga ccgctgggtg ttccccgaga ccaacactgg catcattgt 299

<210> 1407
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 1407

gcccgtccaa gcccttcgcc ggcgctagga tctcggggtc tctccacatg accatocaga 60
ccgcgcctct catcgagacc ctcaccgcgc tcggcgccga ggtccgctgg tgctcctgca 120
acatcttctc cagcgaggac cagcgccgcg ccgccatcgc gcgcnactcg gccgcgctgt 180
tcgcctggna gggggagacc ctgaggagt actggtggtg caccgagcgc tgctctgact 240
ggngcgange gggcggnccc gacctcatcg tcgacgacgg cggcgacgcc acgtgctc 299

<210> 1408
<211> 303
<212> nucleic acid
<213> Zea mays

<400> 1408

cagcgaggac cagcgccgcg ccgccatcgc gcgcgactcg gccgcgctgt tcgcctggna 60
gggggagacc cttgaggagt actggtggtg caccgagcgc tgctctgact ggggcgaggg 120
ggggggcccc gacctcatcg tcgacgacgg cggcgacgcc acgtgctca tccacgaggg 180
tgtcaaggcc gaggaggagt acgagaagac cggcaagatc cccgacccgg agtccaccga 240
caacgctgag ttcaagatcg tgctcaccat catccgcgac gggctcaagg ctgaccccaa 300
gaa 303

<210> 1409
<211> 494
<212> nucleic acid
<213> Zea mays

<400> 1409

gggttngnga aattcctacg ntctcggnat cggtcntnaa ttcacgggtc gacccacgct 60

ccanttaaa ttccncatct cccatgattc aatttcgcga agttctccct cctctgcccc 120
atggcgctct ctgtgganaa gacctcgtct ggacgggagt acaagggtcaa ggatctccgc 180
angcggactt cggncgcctc gagattgagc tggccgaggt cgaaatgccc ggctcttggc 240
gttgccgcgc cgagttcggc ccgtcnaagc ccttcgctgg cgctaggatc tcgggtctct 300
ccacatgacc atccaaaccg ccgtcctcat cgagaccctc accgcgctcg gcgcgaggtc 360
cgctggtgct cctgcaacat cttctccacg cagaccacg ccgccgcgc catgcgcgcg 420
actcgccgcg cgntgttcnc cctggaangg gggaaaacct ccaagaanta ctgtggttca 480
ancgagccgc tgnt 494

<210> 1410
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 1410
cnggggtctcn ccacatgacc atccagaccg ccgtcctcat cgagaccctc accgcgctcg 60
gcgccgaggt ccgcnggtgc nctgcaaca ncttcnccac gcaggaccac gccgcngccg 120
ccatcgcgcg cgantcggcc gcngtgtncg cctggaaggg ggagaccctc gangagtact 180
ggnggtgcac cgagcgctgc ctcgactggn gcgangcggg cggccccgac ctcatcgctg 240
acgacggcgg cgacgccacg ctgctcatcc acgagggtgt caangccgag gaggattac 299

<210> 1411
<211> 302
<212> nucleic acid
<213> Zea mays

<400> 1411
cgtcgacnac ggcggcgacg ccacgctgct catccacgag ggtgtcaagg ccgangagga 60
ttacgagaag accggcaaga tctccgaccc ggagtccacc gacaacgctg agttcaagat 120
cgtgctcacc atcatccgcg acgggctcaa ggctgacccc aagaagtacc gcaagatgaa 180
ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 240
gcagganacc ggcgcctctc tcttcctgac cattaacgtc nacgattccg tcaccaagag 300
ca 302

tctgaggaga ccaccacggg tgtcaagagg ctctaccaga tgcaggagac cggcgcacctc 60
ctcttccctg ccattaacgt caacgattcc gtcaccaaga gcaagtttga caacctgtat 120
ggttgccgcc actcgctccc tganggtctg atgagggcca ctgacgttat gatcgccgga 180
aaggttgcn g tggntgcn g atacggtgat gtcggcaagg gttgtgctgc tgccctcaag 240
caggctggtg cccgtgtcat tgtgaccgan atnacccca tctgtgctc caggctctga 300
tggaggggtct t 311

<210> 1415
<211> 280
<212> nucleic acid
<213> Zea mays

<400> 1415

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cgccgcgcgc atcgcgcgcg actcggcgcg cgtgttcgcc tggaaggggg agacctcga 120
ggagtaactgg tgggtgcaccg agcgctgect cgactggggc gaggcgggcg gccccgacct 180
catcgctgac gacggcgggc acgccacgct gctcatccac gagggtgtca aggccgagga 240
ggattacgag aagaccggca agatccccga cccggagtcc 280

<210> 1416
<211> 295
<212> nucleic acid
<213> Zea mays

<400> 1416

gagatcgacc ccattctgcgc cctccagget ctgatggagg gtcttcaggt ccttcccttg 60
gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 120
atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tggccacttt 180
gacaatgaaa ttgatatgct cggccttgag acctaccctg gcgtcaagcg catcaccatc 240
aagccccaga ctgaccgctg ggtgttcccc gagaccaaca ctggcatcat tgtcc 295

<210> 1417
<211> 349
<212> nucleic acid
<213> Zea mays

<400> 1417

ctggcgtcaa ggcgcatcacc atcaagcncc agactgaccg ctgggtgttc cccgagacca 60
aactggcat cattgtcctt gctgaggtc gcctgatgaa ccttgggtgt gctactggcc 120
atcctagctt tgtcatgtcc tgetcattca ctaaccaggt cattgcccc attgaactgt 180
ggaaggagaa gagctctggc aagtatgaga agaaggtgta tgtgtcccc aagcaccttg 240
atgagaaggt tgctgctctc cacttgggca agcttgggtc caagctggac caagctcacc 300
aagtctcagg ccgatacatc agcgtgccga tcgnggtcct acaagcctg 349

<210> 1418

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 1418

ctgatgaggg ccaccgacgt tatgatcgcc ggtaagggtg ccgtggtctg cggatacggg 60
gatgttggca agggtttgtc cgtgcactc aagcaggctg gtgcccggtg catttgtgacc 120
gagatcgacc ccattctgcgc cctccaggct ctgatggagg gtcttcagggt ccttcccttg 180
gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 240
atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tg 292

<210> 1419

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1419

agaggctcta ccagatgcag gagaccggcg cctcctctt cctgccatt aacgtcaacg 60
attccgtcac caagagcaag ttgacaacc tgtatgggtg ccgccactca ctccctgatg 120
gtctgatgag ggccaccgac gttatgatcg ccgtaagggt tgccgtggtc tgcggatacg 180
gtgatgttgg caagggttgt gccgtgcac tcaagcaggc tgggtgccgt gtcatttgtga 240
ccgagatcga ccccatctgc gccctccagg ctctgatgga gggctctt 287

<210> 1420

<211> 304

<212> nucleic acid

<213> Zea mays

<400> 1420

gtgcnctcca ngctctgatg gaggggtcttc aggtccttcn cttggangac gttgtctctg 60
aagctgacat ctctgtgacc accactggca acaangatat catcatgggt gaccacatga 120
ggaagatgaa gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata 180
tgctcggcct tgagacctac cctggcgctca agcgcatcac catcaagccc cagactgacc 240
gctgggtggt ccccgagacc aacactggca tcatgtcttg ctganggtcg cctgatgaac 300
cttg 304

<210> 1421

<211> 283

<212> nucleic acid

<213> Zea mays

<400> 1421

ggagtactgg tgggtgcaccg agcgctgcct cgactggggc gaggcgggcg gccccgacct 60
catcgtcgac gacggcgggcg acgccacgct gtcctccac gaggggtgtca aggccgagga 120
ggattacnag aagaccggca agatccccga cccggagtcc accgacaacg ctgagttcaa 180
gacgtgtctc accatcatcc gcgacgggct caaggctgac cccaagaagt accgcaagat 240
gaaggagagg cttgtcggcg tctctgagga gaccaccacg ggt 283

<210> 1422

<211> 420

<212> nucleic acid

<213> Zea mays

<400> 1422

ggatatgctc cgggcttgag acctaccctg gcgtaagcg catnaccatc aagccccaga 60
ctgaccgctg ggtgttcccc gagaccaaca ctggcatcat tgncttgct gagggctgcc 120
tgatgaacct tgggtgtgct actggccatn ctagctttgt catgtctgc tcattcacta 180
accaggtcat tgcccaactt gaactgtgga aggagaagag ctctggcaag tatnanaaga 240
angtgtatgt gctncccaag caccttgatn agaangntgn tgntctncac ttgggcaagc 300
ttggtgcaa nctnaccaag cttaccaaag tcttaagncc gctacnttaa cctgcccntc 360

gaaggttcct tccaacctnn ccacnaaccg gtcttagnaa gcnnnacaac ggttngaant 420

<210> 1423
 <211> 311
 <212> nucleic acid
 <213> Zea mays

<400> 1423

gaccggcgcc ctctcttccc ctgccattaa cgtcaacgat tccntcacca agancaagtt 60

tgacaacctg tatggttnen gccactcgct cctgatggt ctgatgagg ccactgacgt 120

tatgatcgcc ggaaagggtg cctggtctg cngatacgt gatgtcggca aggnttggtg 180

tgctgccctc aagcaggctg gtgcccggtg cattgtgacc ganatcgacc ccactcttcc 240

cctccaggct ctgatggagg gtcttcagg ccttccttg gaagacgttg tctctgaagc 300

tgacatcttc g 311

<210> 1424
 <211> 283
 <212> nucleic acid
 <213> Zea mays

<400> 1424

cgacgccacg ctgctcatcc acgagggtgt caaggccgag gaggagtacg agaagaccgg 60

caagatcccc gaccgggagt ccaccgacaa cgctgagttc aagatcgtgc tcaccatcat 120

ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga ggcttgctcg 180

cgtctctgag gagaccacca cgggtgtcaa gaggctctac cagatgcagg agaccggcgc 240

cctctcttc cctgccatta acgtcaacga ttccgtcacc aag 283

<210> 1425
 <211> 369
 <212> nucleic acid
 <213> Zea mays

<400> 1425

aattcctca cnaaaagcaa gtttganaac ctgtatggtt gccgcnann actccctgat 60

ggnetgatna gggccaccga cgttatgatc gccgtaagg ttgccgtggn ctgcngatac 120

cgtgatgttg gcaanggttg tgccnctgca ctcaagcagg ctgntgcccg tgtcattgtg 180

accgagatcg annccatctg cgccctccac gctctgatgg atgggtcttc aagtccttcc 240
 cttggaggac gttgtctcgg gaagctgaca tcttcgtgac caccactggc aacaaggata 300
 tcatcaatgg gttgancaca tgaaggaacg atgaaggaca atggcantgt ctgcaacatt 360
 gggcaacnt 369

<210> 1426
 <211> 278
 <212> nucleic acid
 <213> Zea mays
 <400> 1426

gcaagatccc cgaccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 60
 tccgcgacgg gctcaagggt gacccaaga agtaccgcaa gatgaaggag aggcttgtcg 120
 gcgtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 180
 ccctcctctt ccttgccatt aacgtcaacg attccgtcac caagagcaag tttgacaacc 240
 tgtatgggtg ccgccactcg ctcctgatg gtctgatg 278

<210> 1427
 <211> 275
 <212> nucleic acid
 <213> Zea mays
 <400> 1427

cttcgtgacc accactggca acaaggatat catcatgggt gaccacatga ggaagatgaa 60
 gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct 120
 tgagacctac cctggcgtca agcgcatcac catcaagccc cagactgacc gctgggtggt 180
 ccccgagacc aacactggca tcattgtcct tgctgagggt cgcctgatga accttgggtg 240
 tgctactggc catcctagct ttgtcatgtc ctgct 275

<210> 1428
 <211> 275
 <212> nucleic acid
 <213> Zea mays
 <400> 1428

tgacatcttc gtgaccacca ctggcaacaa ggatatcatc atggttgacc acatgaggaa 60

gatgaagaac aatgccattg tctgcaacat tggccacttt gacaatgaaa ttgatatgct 120
 cggccttgag acctaccctg gcgtaagcg catcaccatc aagccccaga ctgaccgctg 180
 ggtgttcccc gagaccaaca ctggcatcat tgtccttgct gagggtcgcc tgatgaacct 240
 tgggtgtgct actggccatc ctagctttgt catgt 275

<210> 1429
 <211> 294
 <212> nucleic acid
 <213> Zea mays
 <400> 1429

caccatcaag cccagactg accgctgggt gttccccgag accaactctg gcatcattgt 60
 ccttgctgag ggctgcctga tgaaccttg gtgtgctact ggccatccta gctttgtcat 120
 gtctgctca ttactaacc aggtcattgc ccaacttgaa ctgtggaagg agaagagctc 180
 tggcaagtat gagaagaagg tgtatgtgct cccaagcac cttgatgaga aggttgctgc 240
 tctccacttg ggcaagcttg gtgccaagct gaccaagctc accaagtctc aggc 294

<210> 1430
 <211> 276
 <212> nucleic acid
 <213> Zea mays
 <400> 1430

gctgacatct tcgtgaccac cactggcaac aaggatatca tcatggttga ccacatgagg 60
 aagatgaaga acaatgccat tgtctgcaac attggccact ttgacaatga aattgatatg 120
 ctggccttg agacctacc ttggctcaag cgcaccacca tcaagcccca gactgaccgc 180
 tgggtgttcc ccgagaccaa cactggcatc attgtccttg ctgagggtcg cctgatgaac 240
 cttgggtgtg ctactggcca tcctagcttt gtcatg 276

<210> 1431
 <211> 288
 <212> nucleic acid
 <213> Zea mays
 <400> 1431

ctcatggcgt gccgcgccga gttcgggccc tccaagccct tcgccggcgc taggatctcg 60

gggtctctcc acatgaccat ccagaccgcc gtctctcatcg agaccctcac cgcgctcggc 120
gccgaggtcc gctggtgctc ctgcaacatc ttctccacgc aggaccacgc cgccgccgcc 180
atcgcgcgcg actcgccgc cgtgttcgcc tggnaagggg agacccttga ggagtactgg 240
tggtgcaccg agcgctgcct tgactggggc gangcggggc gccccgac 288

<210> 1432
<211> 285
<212> nucleic acid
<213> Zea mays

<400> 1432

tgcaggagac cggcgccctc ctcttccctg ccattaacgt caacgattcc gtcaccaaga 60
gcaagtttga caacctgtat ggttgccgcc actcgctccc tgatggtctg atgagggcca 120
ctgacgttat gatcgccgga aaggttgccg tggctctgcg atacggtgat gtcggcaagg 180
gttgtgctgc tgccctcaag caggctggtg cccgtgtcat tgtgaccgag atcgacccca 240
tctgtgccct ccaggctctg atggagggtc ttcaggctct tccct 285

<210> 1433
<211> 280
<212> nucleic acid
<213> Zea mays

<400> 1433

atcgacccca tctgtgccct ccaggctctg atggagggtc ttcaggctct tcccttgag 60
gacgttgtct ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg 120
gttgaccaca tgaggaagat gaagaacaat gccattgtct gcaacattgg ccactttgac 180
aatgaaattg atatgctcgg ccttgagacc taccctggcg tcaagcgcat caccatcaag 240
ccccagactg accgctgggt gttccccgag accaacactg 280

<210> 1434
<211> 316
<212> nucleic acid
<213> Zea mays

<400> 1434

cgaccccatc tgcncctcc aggtctctgat ggagggtctt caggctcttc ccttgaggga 60

cgttgtctcg gaagctgaca tcttcgtgac caccactggc aacaaggata tcatcatggt 120
 tgaccacatg aggaagatga agaacaatgc catttgtctgc aacattggcc actttgacaa 180
 tgaaattgat atgctcggcc ttgagacctc ccttggcgtc aagcgcatca ccatcaagcc 240
 ccagactgac cgctgggtgt tccccgagac caacactggc atcatgtcct tgctgaaggt 300
 cgctgatga acttgg 316

<210> 1435
 <211> 298
 <212> nucleic acid
 <213> Zea mays

<400> 1435

gcccgccctc atggcgtgcc gcgcgagtt cggcccgctc aagcccntcg ccggcgctag 60
 gatctcgggg tototccaca tgaccatcca gaccgcgctc ctcctcgaga ccctcaccgc 120
 gctcggcgcc gaggtccgct ggtgctcctg caacatcttc tccacgcagg accacgccgc 180
 cgccgccatc gcgcgcgact cggccgccgt gttcgctcgg aaagggggag accctcgagg 240
 agtactggtg gtgcaccgag cgctgctcga ctggggcgaa gcggggcgcc cgacctca 298

<210> 1436
 <211> 299
 <212> nucleic acid
 <213> Zea mays

<400> 1436

agaagaccgg caagatcccc gaccgggagt ccaccgacaa cgctgagttc aagatcgtgc 60
 tcaccatcat ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga 120
 ggcttgctcg cgtctctgag gagaccacca cgggtgtcaa gaggtcttac cagatgcagg 180
 agaccggcgc cctcctcttc cctgccatta acgtcaacga ttccgtcacc aagagcaagt 240
 ttgacaactg tatggttgcc gcanttcgtt ccttgatggt ttgatgaggg cactgang 299

<210> 1437
 <211> 279
 <212> nucleic acid
 <213> Zea mays

<400> 1437

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aacgctgagt tcaagatcgt gctcaccatc atccgcgacg ggctcaaggc tgaccccaag 120
aagtaccgca agatgaagga gagggcttgc gggcgctctc gagggagacca ccacgggtgt 180
caagaggctc taccagatgc aggagaccgg cgcctctctc ttccttgcca ttaacgtcaa 240
cgattccgtc accaagagca agtttgacaa cctgtatgg 279

<210> 1438
<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1438

gcaagatccc cgacccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 60
tccgcgacgg gctcaaggct gaccccaaga agtaccgcaa gatgaaggag aggcttgtcg 120
gcgctctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 180
ccctcctctt ccttgccatt aacgtcaacg attccgtcac caagagcaag ttgacaact 240
gtatggttgc cgccactgc tccctgatgg tctgatg 277

<210> 1439
<211> 318
<212> nucleic acid
<213> Zea mays

<400> 1439

atttccccat ctcccagatc caattcgca gttctcctc ctctgccgcc atggcgctct 60
ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg caggcggact 120
tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg gcgtgccgcg 180
ccgagttcgg cccgtccaag cccttcgccg gcgctaggat ctcggggtct ctccacatga 240
ccatccagac cgcgctctc atcgagacc tcaccgcgt cggcgccgag gtccgctggt 300
gtcctgcaa catcttct 318

<210> 1440
<211> 249
<212> nucleic acid
<213> Zea mays

<400> 1440

cacatgacca tccagaccgc cgtcctcatc gagaccctca ccgcgctcgg cgccgaggtc 60

cgctggtgct cctgcaacat cttctccacg caggaccacg ccgccgccgc catcgcgcg 120

gactcggccg ccgtgttcgc ctggaagggg gagaccctcg aggagtactg gtggtgcacc 180

gagcgtgcc tcgactgggg cgaggcgggc ggccccgacc tcatcgtcga cgacggcggc 240

gacgccacg 249

<210> 1441

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1441

cccagatcca attcgcgagt nonccctcct ctgcggccat ggcgctctct gtggagaaga 60

cctcgtctgg acgggagtac aagggtcaagg atctctcgca ggcggacttc ggccgcctcg 120

agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gaggttcgcc 180

cgtccaagcc cttcgccggc gctaggatct cggggctctct ccacatgacc atccagaccg 240

ccgtcctcat cgagaccctn accgcgctcg gcgccgaggt ccgctggtgc tcctgcaaca 300

tcttctcca 309

<210> 1442

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1442

gtacgagaag accggcaaga tccccgaccc ggagtccacc gacaacgctg agttcaagat 60

cgtgctcacc atcatccgcg acggggtcaa ggctgacccc aagaagtacc gcaagatgaa 120

ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 180

gcaggagacc ggccgccctcc tcttcctgc cattaacgtc aacgattccg tcaccaagag 240

caagtttgac aacctgtatg gttgccgcca ctact 276

<210> 1443

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1443

gtgaccgana tcgaccccat ctgtncctc caggctctga tggaggggtct tcaggtcctt 60
cccttgaggg acgttgtctc tgaagctgac atnttcgtga ccaccactgg caacaaggat 120
atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 180
cactntgaca atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcatc 240
accatcaagc cccagactga cgcgtgggtg ttcccc 276

<210> 1444

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 1444

agaagaccgg caagatcccc gaccgggagt ccaccgacaa cgctgagttc aagatcgtgc 60
tcaccatcat ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga 120
ggcttgctgg cgtctctgag gagaccacca cgggtgtcaa gaggtctctac cagatgcagg 180
agaccggcgc cctcctcttc cctgccatta acgtcaacga ttccgtcacc aagagcaagt 240
ttgacaacct gtatggttgc cgccactcgc 270

<210> 1445

<211> 261

<212> nucleic acid

<213> Zea mays

<400> 1445

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catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 120
tgacaatgaa attgatatgc tcggccttga gacctacctt ggcgtaagc gcatcaccat 180
caagccccag actgaccgct ggggtgtccc cgagaccaac actggcatca ttgtccttgc 240
tgagggtcgc ctgatgaacc t 261

<210> 1446

<211> 291

<212> nucleic acid

<213> Zea mays

<400> 1446

tntgacaacc tgtatggttg cggccactcg ctccctgatg gtctgatgag ggccactgac 60
 gttatgatcg ccggaaggt tgccgtggtc annccgatac ggtgatgtcg gcaagggttg 120
 tgctgctgcc ctcaagcagg ctggtgcccg tgtcatttg accgagatcg accccatctg 180
 tgccctccag gctctgatgg agggctctca ggtccttccc ttggaggacg ttgtctctga 240
 agctgacatc ttcgtgacca ccactggcaa caaggatatc atcatggttg a 291

<210> 1447

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 1447

actggggcga gggggggcgc cccgacctca tcgtcgacga cggcggcgac gccacgctgc 60
 tcatccacga ggggtgtcaag gccgaggagg agtacgagaa gaccggcaag atccccgacc 120
 cggagtccac cgacaacgct gagttcaaga tcgtgctcac catcatccgc gacgggctca 180
 aggctgaccc caagaagtac cgcaagatga aggagagctt gtcggcgtct taaggagacc 240
 accaggggtgt caagaagctc taccagatgc aagaaaccgg cgccctcctc ttccctgcc 300
 ttaacgtnac gatccg 316

<210> 1448

<211> 273

<212> nucleic acid

<213> Zea mays

<400> 1448

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 gccgaggagg agtacgagaa gaccggcaag atccccgacc cggagtccac cgacaacgct 120
 gagttcaaga tcgtgctcac catcatccgc gacgggctca aggctgaccc caagaagtac 180
 cgcaagatga aggagaggct tgctggcgctc tctgaggaga ccaccacggg tgncaagag 240
 gctctaccag atgcaggaga ccggcgccct cct 273

<210> 1449

<211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 1449

 cngacgctgg gtggctcctg caacatcttc tccacgcagg accacgccgc cgccgccatc 60
 gcgcgcgaca cggccgccgt gttcgccctgg aagggggaga ccctcgagga gtactgggtg 120
 tgcaccgagc gctgcctcga ctggggcgag gcggggcgcc ccgacctcat cgtcgacgac 180
 ggcggcgacg ccacgctgct catccacgag ggtgtcaagg ccgaggagga ttacgagaag 240
 accggcaaga tccccgacct ggagtccacc g 271

<210> 1450
 <211> 275
 <212> nucleic acid
 <213> Zea mays

 <400> 1450

 aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca ctgctccct 60
 gatggtctga tgagggccac tgacgttatg atcgccggaa aggttgccgt ggtctgcgga 120
 tacggtgatg tcggcaaggg ttgtgctgct gccctcaagc aggctggtgc ccgtgtcatt 180
 gtgaccgaga tcgaccccat ctgtgccttc caggctctga tggagggttn caaggctcctn 240
 cccttgaggg acgttgtctc ngaagatgac atctt 275

<210> 1451
 <211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 1451

 gagatcgacc ccattctgcgc cctccaggct ctgatggagg gtcttcaggc ccttcccttg 60
 gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 120
 atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tggccacttt 180
 gacaatgaaa ttgatatgct cggccttgag acctaccctg gcgtcaagcg catcaccatc 240
 aagccccaga ctgaccgctg ggtgttcccc g 271

<210> 1452

<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1452

cattaacgtc aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca 60
ctcactccct gatggtctga tgagggccac cgacgttatg atcgccggta aggttgccgt 120
ggtctgcgga tacggtgatg ttggcaaggg ttgtgccgt gcaactcaagc aggctggtgc 180
ccgtgtcatt gtgaccgaga tcgaccccat ctgcgccctc caggctctga tggagggctct 240
tcaggtcctt cccttgaggg acgttgtctc ggaactg 277

<210> 1453
<211> 273
<212> nucleic acid
<213> Zea mays

<400> 1453

ctaccctggc gtcaagcgca tcaccatcaa gcccagact gaccgctggg tgttccccga 60
gaccaacact ggcattcattg tccttgctga gggtcgcctg atgaaccttg ggtgtgctac 120
tggccatcct agctttgtca tgtcctgctc attcactaac caggtcattg cccaacttga 180
actgtggaag gagaagagct ctggcaagta tgagaagaag gtgtatgtgc tccccaaagca 240
ccttgatgag aagggttgctg ctctccactt ggg 273

<210> 1454
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 1454

ggaggggtctt caggtccttc ccttgaggga cgttgtctct gaagctgaca tcttcgtgac 60
caccactggc aacaaggata tcatcatggt tgaccacatg aggaagatga agaacaatgc 120
natgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct tgagacctac 180
cctggcgta agcgcatac catcaagccc cagactgacc gctgggtgtt ccccgagacc 240
aacactggca tcattgtcct tgctgagggt cgctgatgaa cttgggtgtg tatggccac 299

<210> 1455

<211> 282
 <212> nucleic acid
 <213> Zea mays

<400> 1455

ccgccactcg ctccctgatg gtctgatgag ggccactgac gttatgatcg ccggaaaggt 60
 tgccgtgggc tgcggatacg gtgatgtcnn gcaaggggtg tgctgctgcc ctcaagcagg 120
 ctggtgcccc tgtcattgtg accgagatcg accccatctg tgccctccag gctctgatgg 180
 agggctttca ggtccttccc ttggaggacg ttgtctctga agctgacatc ttcgtgacca 240
 ccaactggcaa caaggatata atcatgggtg accacatgag ga 282

<210> 1456
 <211> 297
 <212> nucleic acid
 <213> Zea mays

<400> 1456

ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct ctgaggagac 60
 caccacgggt gtcaagagggc tctaccagat gcaggagacc ggcgcctcc tcttccctgc 120
 cattaacgtc aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca 180
 ctgctccct gatggtctga tganggccac tgacgttatg attcgccgga aagggtgccg 240
 tgggtctgogg atacggtgat gtcggcaang gttgtgtgct gccctcaagc angctgg 297

<210> 1457
 <211> 130
 <212> nucleic acid
 <213> Zea mays

<400> 1457

gngatcgacc ccattctgcgc cctccagnct ctnatggagg gtcttcaggt ccttcccttg 60
 naggacgttg tctcggaagc tgacatcttc ggtgaccacc actggcaaca aggatatan 120
 ncatgggttg 130

<210> 1458
 <211> 304
 <212> nucleic acid
 <213> Zea mays

<400> 1458

catctcccag atccaattcg cgagttctcc ctctctgcg gccatggcgc tctctgtgga 60
gaagacctcg tctggacggg agtacaaggt caaggatctc tcgcaggcgg acttcggccg 120
cctcgagatt gagctggcgg aggtcgaaat gcccggcctc atggcgtgcc gcgccgagtt 180
cgccccgtcc aagcccttcg cgggcgctag gatctcgggg tctctccaca tgaccatcca 240
gaccgcgctc ctcatcgaga cctcaccgc gctcggcgcc gaggtccgct ggtgctcctg 300
caac 304

<210> 1459

<211> 512

<212> nucleic acid

<213> Zea mays

<400> 1459

gnggnnnng agtgnntnnt aatttgagg naaaatgtaa nggnaatctt cgtaccggtc 60
cggaannntc gaccacgcg tccgccacg cgtccggacc aacactggca tcattgtcct 120
tgctgagggt cgnctgatga ncttggggtg tgctactggc catcctagct ttgncatgtc 180
ctgtcatctc actaaccagg tcattgcca acttgaactg tggaaggaga agagctctgg 240
caagtatgaa aagaagggtg atgtntccc caagcacctt gatgagaagg ttgctgctct 300
ccaattgggc aancttggtg ccaagctgac caagctnacc aagtctcagg ccgactacat 360
cagcgtgccg atcgagggtc cctacaagcc tgccactacc ggtactaggc agccagcaca 420
cggnttgcaa ctactcggg cgtgtgtgc tatnaagccg ctactggcc tgnagntatc 480
tnnccgnannc tatggcataa acatanacgg ga 512

<210> 1460

<211> 263

<212> nucleic acid

<213> Zea mays

<400> 1460

gccacgcagg accacgccgc cgccgccatc gcgcgcgact cggccgccgt gttcgctgn 60
aanggggaga cccttgagga gtactgggtg tgnaccgagc gctgccttga ctggggcgag 120
gcgggcggcc ccgacctcat cgtcgacgac ggccgacgac ccacgtgct catccacgag 180

ggtgtcaagg ccgaggagga gtacgagaag accggcaaga tccccgacct ggagtccacc 240
gacaacgctg agttcaagat cgt 263

<210> 1461
<211> 247
<212> nucleic acid
<213> Zea mays

<400> 1461

ggtctctcca catgaaccatc cagaccgccc tctcatcgga gacctcacc gcgctcggcg 60
ccgaggtcgg ctgggtgctcc tgcaacatct tctccacgca ggaccacgcc gccgcccgcca 120
tcgcgcgcga ctgggcccgc gtgttcgcct ggaaggggga gaccttgag gagtactggt 180
ggtgcaccga gcgctgcctt gactggggcg aggcggggcg ccccgacctc atcgtcgacg 240
acggcgg 247

<210> 1462
<211> 260
<212> nucleic acid
<213> Zea mays

<400> 1462

ggaagatgaa gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata 60
tgctcggcct tgagacctac cctggcgctca agcgcatcac catcaagccc cagactgacc 120
gtcgggtggt ccccgagacc aacctggca tcattgtcct tgctgagggc cgctgatga 180
accttgggtg tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg 240
tcattgccc acttgaactg 260

<210> 1463
<211> 272
<212> nucleic acid
<213> Zea mays

<400> 1463

ggccaccgac gttatgatcg ccggttaagg tgccgtggc tcgagatagc gtgatgttgg 60
caagggttgt gccgctgcac tcaagcaggc tgggtcccgt gtcattgtga ccgagatcga 120
cccatctgc gccctccagg ctctgatgga gggctctcag gtccttcctc tggaggacgt 180

tgtctcgga gctgacatct tcttgaccac cactggcaac aaggatatca tcanggttga 240
ccacatgagg aagatganga acaatgccat tg 272

<210> 1464
<211> 253
<212> nucleic acid
<213> Zea mays

<400> 1464

ggttgaccac atgaggaaga tgaagaacaa tgccattgtc tgcaacattg gccactttga 60
caatgaaatt gatatgctcg gccttgagac ctaccctggc gtcaagcgca tcaccatcaa 120
gccccagact gaccgctggg tgttccccga gaccaacact ggcatcattg tccttgctga 180
gggtcgctcg atgaaccttg ggtgtgctac tggccatcct agctttgtca tgtcctgctc 240
attcactaac cag 253

<210> 1465
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 1465

ccacgctgct catccacgag ggtgtcaagg ccgaggagga gtacgagaag accggcaaga 60
tccccgaccc ggagtccacc gacaacgctg agttcaagat cgtgctcacc atcatccgcg 120
acgggctcaa ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct 180
ctgaggagac caccacgggt gtcaagaggc tctaccagat gcaggagacc ggcgccctcc 240
tcttccctgc cattaacgtc a 261

<210> 1466
<211> 261
<212> nucleic acid
<213> Zea mays

<400> 1466

cggggtctct ccacatgacc atccagaccg ccgtctcat cgagaccctc accgcgctcg 60
gcgccgaggt ccgctggtgc tcttgaaca ttttctccac gcaggaccac gccgccgccc 120
ccatcgcgcg cgactcggcc gccgtgttcg cctggaaggg gagacccttg aggagtactg 180

gtggtgcacc gagcgtgcc ttgactgggg cgaggcgggc ggccccgacc tcctcgtcga 240
cgacggcggc gacgcacgt g 261

<210> 1467
<211> 323
<212> nucleic acid
<213> Zea mays

<400> 1467

ctcccgttcc atttccccat ctcccagatc caattcgcca gttctccctc ctctgccgcc 60
atggcgtctt ctgtggagaa gacctgtctt ggacgggagt acaagggtcaa ggatctctcg 120
caggcggact tcggccgctt cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
gcgtgccgcy ccgagttcgg ccggtccaag cccttcgccg ccgctaggat ctcggggtct 240
ctccacatga ccatccagac cgcgtctctc atcgagacct tcaccgcgtt cggcgccgag 300
gtccgcaggt gtcctgcaa cat 323

<210> 1468
<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1468

gtcggcctt gnagacctac cctggcgtca agcgcctcac catcaagccc cagactgacc 60
gtcgggtgtt ccccgagacc aacctggca tcattgtctt tgctgagggt cgctgatga 120
accttgggtg tgctactggc cactctagct ttgtcatgtc ctgctcattc actaaccagg 180
tcattgcca acttgaactg tggaaggaga agagctctgg caagtatgag aagaagggtg 240
atgtgctccc caagcacctt gatgagaagg ttgctgc 277

<210> 1469
<211> 257
<212> nucleic acid
<213> Zea mays

<400> 1469

caaggatatc atcatggttg accacatgag gaagatgaag aacaatgcca ttgtctgcaa 60
cattggccac ttggaacaat gaaattgata tgctcggcct tgagacctac cctggcgtca 120

agcgcatcac catcaagccc cagactgacc gctgggtgtt ccccgagacc aacactggca 180
tcattgtcct tgctgagggg cgctgatga accttgggtg tgctactggc catcctagct 240
ttgtcatgtc ctgctca 257

<210> 1470
<211> 262
<212> nucleic acid
<213> Zea mays

<400> 1470

gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat gcaggagacc 60
ggcgccctcc tcttccctgg ccattaacgt caacgattcc gtcaccaaga gcaagtttga 120
caacctgtat ggttgccgcc actcgtccc tgatgggtctg atgagggcca ctgacgttat 180
gatcgccgga aagggttgcg tgggtctgagg atacgggtgat gtcggcaagg gttgtgctgc 240
tgcaactcaag caggctgggtg cc 262

<210> 1471
<211> 317
<212> nucleic acid
<213> Zea mays

<400> 1471

tatggttgcc gccactcact ccctgatggt ctgatgaggg ccaccgacgt tatgatcgcc 60
ggtaagggttg ccgtgggtctg cggatacggg gatgttgcca agggttgtgc cgctgcactc 120
aagcaggctg gtgcccgtgt cattgtgacc gagatcgacc ccactctgnc cctccaggct 180
ctgatggagg gtcttcagggt ccttcccttg gaggnntngt cacggaagct nanatttctg 240
gaccaccact ggnaacaagg atatcatcat ggttgaccac atgaggaaga tgaanaacat 300
gccattgtct cnaattg 317

<210> 1472
<211> 268
<212> nucleic acid
<213> Zea mays

<400> 1472

cgactggggc gagggcgggc gccccgacct catcgctgac gacggcggcg acgccaacgc 60

tgctcatcca cgagggtgtc aaggccgagg aggattacga gaagaccggc aagatccccg 120
 acccgagtc caccgacaac gctgagttca agatcgtgct caccatcatc cgcgacgggc 180
 tcaaggctga cccaagaag taccgcaaga tgaaggagag gcttgtcggc gtctctgagg 240
 agaccaccac ggggtgtcaag aggtctta 268

<210> 1473
 <211> 274
 <212> nucleic acid
 <213> Zea mays
 <400> 1473

ggcaagggtt gtgtgtgtgc cctcaagcag gctgggtgcc gtgtcattgt gaccgagatc 60
 gaccccatct gtgccctcca ggctctgatg gaggggtcttc aggtccttcc cttggaggac 120
 gttgtctctg aagctgacat cttcgtgacc accactggca acaaggatat catcatggtt 180
 gaccacatga ggaagatgaa gaacaatgcc attgtctgca acattggcca ttgacaatg 240
 aaattgatat gctcggcctt gagacctacc ctgg 274

<210> 1474
 <211> 290
 <212> nucleic acid
 <213> Zea mays
 <400> 1474

gttgtgtgtc tgccctcaag caggctggtg cccgtgtcat tgtgaccgag atcgacccca 60
 tctgtgccct ccaggctctg atggagggtc ttcaggctct tcccttgagg gacgttgtct 120
 ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg gttgaccaca 180
 tgaggaagat gaagaacaat gccattgtct gcaacattgg ccactttgac aatgaaattg 240
 atatgctcgg ccttgagact acctggcgtc aagcgcata catcaagccc 290

<210> 1475
 <211> 300
 <212> nucleic acid
 <213> Zea mays
 <400> 1475

cgagggtgtc aaggccgagg aggagtacga gaagaccggc aagatccccg acccgagtc 60

caccgacaac gctgagttca agatcgtgct caccatcatc cgcgacgggc tcaaggctga 120
 cccaagaag taccgcaaga tgaaggagag gcttgctggc gtctctgagg agaccaccac 180
 ggggtgtcaag aggctctacc agatgcagga gaccggcgcc ctctcttcc ctgccattaa 240
 cgtcaacgat tcgtcaccag agcaagtttg acnactgtat ggttgccgca attcattccc 300

<210> 1476
 <211> 260
 <212> nucleic acid
 <213> Zea mays

<400> 1476

anggtcttca ggtccttccc ttggaggacg ttgtctcgga agctgacatc ttcgtgacca 60
 ccactggcaa caaggatata atcatgggtg accacatgag gaagatgaag aacaatgcc 120
 ttgtctgcaa cattggccac ttgacaatg aaattgatat gtcggcctt gagacctacc 180
 ctggcgtcaa ggcacatcac atcaagcccc agactgaccg ctgggtgttc cccgagacca 240
 aactggcat cattgtcctt 260

<210> 1477
 <211> 295
 <212> nucleic acid
 <213> Zea mays

<400> 1477

attacnagaa gaccggcaag anccccgacc cggagtcac cgacaacgct gagttcaaga 60
 tcgtgctcac catcatccgc nacgggctca aggctgacct caagaagtac cgcaaganga 120
 aggacgaggc ttgtcggcgt ctctgaggag accaccacgg gtgtcaagag gctctaccag 180
 atgcaggaga ccggcgccct cctcttcct gccattaacg tcaangattc cgtcaccaag 240
 agcaagtttg acaacntgta tggttgccgc caactcggct ccctgatggt ctgat 295

<210> 1478
 <211> 278
 <212> nucleic acid
 <213> Zea mays

<400> 1478

gncgngccg ccacgcgcg cgactcggcc gccgtgttcg cctggaaggg ggagaccctt 60

gaggagtact ggtggtgcac cgagcgctgc cttgactggg gcgaggcggg cggccccgac 120
ctcatcgctcg acgacggcgg cgacgccaac gctgctcatc cacgagggtg tcaaggccga 180
ggaggagtac gagaagaccg gcaagatccc cgacccggag tccaaccgac aacgntgagt 240
tcaagatcgt gcttaccatc attcgggacn ggctcaaa 278

<210> 1479
<211> 287
<212> nucleic acid
<213> Zea mays
<400> 1479

gctgagggtc gcttgatgaa ccttgggtgt gctactggcc atcctagctt tgtcatgtcc 60
tgctcattca ctaaccaggt cattgcccac cttgaactgt ggaaggagaa gagctctggc 120
aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgctgctctc 180
catttgggca agcttgggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 240
agcgtgccga tcgaggggtcc ctacaagcct gccactacc ggtacta 287

<210> 1480
<211> 306
<212> nucleic acid
<213> Zea mays
<400> 1480

cgtcaccaag agcaagtttg acaacctgta tggttgccgc catcgctccc tgatgggtctg 60
aatgagggcc actgacgtta tgatcgccgg aaaggttgcc gtggtctgcy gatacgggtga 120
tntcggcaan ggttgtgctg ctgccctcaa gcaggctggt gcccggtgtca ttgtgaccga 180
gatcgacccc atctgtgccc tccaggctct gatggagggt cttcagggtcc ttcccttgga 240
ggacgttgtc tctgaagctg acatcttcgt gaccaccact ggcaacaagg atatcacatg 300
gttgac 306

<210> 1481
<211> 314
<212> nucleic acid
<213> Zea mays
<400> 1481

ctctgaagct gacatcttcg tgaccaccac tggcaacaag gatatcatca tggttgacca 60
catgaggaag atgaagaaca atgccattgt ctgcaacatt ggccactttg acaatgaaat 120
tgatatgctc ggccttgaga cctaccctgg cgtcaagcgc atcaccatca agccccagac 180
tgaccgctgg gtgttccccg agaccaaacac tggcatcatt gtccttgctg anggtcgcc 240
gatgaacttg ggtgtgtatg gccatcctag tttgtcatgt cctgtcatna ctaaccagtc 300
attgnccaat tgaa 314

<210> 1482
<211> 270
<212> nucleic acid
<213> Zea mays

<400> 1482
atgccattgt ctgcaacatt ggccactttg acaatgaaat tgatatgctc ggccttgaga 60
cctaccctgg cgtcaagcgc atcaccatac aagccccaga ctgaccgctg ggtgttcccc 120
gagaccaaca ctggcatcat tgtccttgct gagggtcgcc tgatgaacct tgggtgtgct 180
actggccatc ctagctttgt catgtcctgc tcattcacta accagggtcat tgcccaactt 240
gaactgtgga aggagaagag ctctggcaag 270

<210> 1483
<211> 266
<212> nucleic acid
<213> Zea mays

<400> 1483
caccgacgtt atgatcgccg gtaagggtgc cgtggtctgc ggatacgggtg atgttgggcaa 60
ggnttgtgcc gctgcactca agcaggctgg tgcccgtntc attgtgaccg agatcgaccc 120
catctgcnn ctcangctc tgatggaggg tcttcaggtc cttcccttgg aggacgttgt 180
ctcggaagct gacatcttcg tgaccaccac tggcaacaag gatatcatca tggttgacca 240
catgaggaag atgaagaaca atgcca 266

<210> 1484
<211> 312
<212> nucleic acid
<213> Zea mays

<400> 1484

ctcgttccat ttccccatct cccagatcca attcgcgagt tctccctcct ctgcggccat 60
 ggcgctctct gtggagaaga cctcgtctgg acgggagtag aaggtcaagg atctctcgca 120
 ggcggaactt ggccgcctcg agattgagct ggccgaggtc gaaatgcccg gctcatggc 180
 gtgccgcgnc gagttcggcc cgtccaagcc cttcgccggc gctaggatct cggggtctct 240
 ccacatgacc atccagaccg cgtctctcat cgagaccctc accgcgctcg gcgccgaggt 300
 ccgctggtgc tc 312

<210> 1485

<211> 271

<212> nucleic acid

<213> Zea mays

<400> 1485

aagccccaga ctgaccgctg ggtgttcccc gagaccaaca ctggcatcat tgtccttgc 60
 gagggtcgcc tgatgaacct tgggtgtgct actggccatc ctagctttgt catgtcctgc 120
 tcattcacta accaggtcat tgcccaactt gaactgtgga aggagaagag ctctggcaag 180
 tatgaaagaa ggtgtatgtg ctccccaagc accttgatga gaaggttgct gctctccact 240
 tgggcaagct tggtgccaag ctgaccaagc t 271

<210> 1486

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1486

actggggcga ggccggcgcc cccgacctca tcgtcgacga cggcgggcgac gccaacgctg 60
 ctcatccacg aggggtgtcaa ggccgaggag gattacgaga agaccggcaa gatccccgac 120
 ccggagtcca ccgacaacgc tgagttcaag atcgtgtcga ccatcatccg cgacgggctc 180
 aaggctgacc ccaagaagta ccgcaagatg aaggagaggc ttgtcggcgt ctctgaggag 240
 accaccacgg gtgtcaagag gtctaccaga tgcag 275

<210> 1487

<211> 407

<212> nucleic acid

<210> 1490
 <211> 303
 <212> nucleic acid
 <213> Zea mays
 <400> 1490
 gagggccacc gacgttatga tcgccggtaa ggttgccgng gtctgcggnt acggtgatgt 60
 tggcaagggt tgtgccgctg cactacaagc aggctgggtc cegtgtcatt gtgaccgagn 120
 atcgacccca tctgcgccct ccaggctctg atggagggtc ttcaggtcct tcccttgagg 180
 gacgttgtct cggaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg 240
 gttgaccaca tgaggaagat gaagaacaat gccatgtctg caacntggcc atttgacang 300
 aat 303

<210> 1491
 <211> 268
 <212> nucleic acid
 <213> Zea mays
 <400> 1491
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 gtacgagaag accggcaaga tccccgaccc ggagtccacc gacaacgctg agttcaagat 120
 cgtgctcacc atcatccgcg acggggtcaa ggctgacccc aagaagtacc gcaagatgaa 180
 ggagangctt gtcggcgtct ttgaggagac caccangggg gtcaagaggt ctaccagatg 240
 caggagaccg gcgccctcct cttccctg 268

<210> 1492
 <211> 278
 <212> nucleic acid
 <213> Zea mays
 <400> 1492
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 agaccggcgc cctcctcttc cctgncatta acgtcaacga ttccgtcacc aagagcaagt 120
 ttgacaacct gtatggttgc cgccactcgc tccctgatgg tctgatgagg gccactgacg 180
 ttatgatcgc cggaaagggt gccgtggtct gcggatacgg tgatgtcggc aagggttgtg 240

ctgctgccct caagcaggct ggtgccgtgt catgtgac 278

<210> 1493
 <211> 282
 <212> nucleic acid
 <213> Zea mays

<400> 1493

gctognnctt gannnetacc cngngtcaa gngcatcnc atcangcccc agactgancg 60
 ctgggnnttc ccnanaacca aactggcat cantgtcctt gctganggtc gcctgatgaa 120
 ccttnggtgt gnaactggcc atcctagctt tgtcangtnc tgctcattca ctaaccaggt 180
 cattgcccac cttgaactgt ggaaggagaa gagctctggc aagtatgaga agaagggtga 240
 tgtgctcccc aagcaccttg atgagaaggt tgctganctc ca 282

<210> 1494
 <211> 305
 <212> nucleic acid
 <213> Zea mays

<400> 1494

gccatggcgc tctctgtgga gaagacctcg tctggacggg agtacaaggt caaggatctc 60
 tcgcaggcgg acttcggccg cctcgagatt gagctggccg aggtcgaaat gcccggcctc 120
 atggcgtgcc gcgcgcaggt cgccccgtcc aagcccttcg ccggcgctag gatctcgggg 180
 tctctccaca tgaccatcca gaccgccgtc ctcatcgaga cctcaccgc gctcggcgcc 240
 gaggtccgtg gtgtcctgca acatttctcc acnaggacca gccgcgcgca tgcgcggaan 300
 ggcg 305

<210> 1495
 <211> 284
 <212> nucleic acid
 <213> Zea mays

<400> 1495

cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60
 cctcgtctgg acgggagtag aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120
 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180

cgccaagcc cttcgccggc gctaggatct cgggggtctct ccacatgacc atccagaccg 240
ccgtcctcat cgagaccctc accgcgctcg gcgccgaggt ccgt 284

<210> 1496
<211> 263
<212> nucleic acid
<213> Zea mays

<400> 1496

ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 60
gcaggagacc ggcgcctcc tcttcctgc cattaacgtc aacgattccg tcaccaagag 120
caagtttgac aacctgtatg gttgccgcca ctgcctcct gatggtctga tgagggccac 180
tgacgttatg atcgccggaa aggttgccgt ggtctgcgga taccgtgatg tcggcaaagg 240
gttgtgctgc tgccctcaag cag 263

<210> 1497
<211> 347
<212> nucleic acid
<213> Zea mays

<400> 1497

ctccctcaact ccggttccat ttccccatct ccagatcca attcgcgagt tctccctcct 60
ctgccgcat ggcgtctct gtggagaaga cctcgtctgg acgggagtag aaggtaagg 120
atctctcgca ggcggaattc ggccgcctcg agattgagct ggccgaggtc gaaatgcccg 180
gcctcatggc gtgccgcgcc gagttcggcc cgtccaagcc cttcgccggc gctaggatct 240
cgggggtctct ccacatgacc atccagaccg ccgtcctcat cgagaccctc accgcgctcg 300
gcgccgaagt ccgtggtgtc tgcaacatct tctccacgan gaccacg 347

<210> 1498
<211> 275
<212> nucleic acid
<213> Zea mays

<400> 1498

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gtcattcac nnaccaggtc attgccaac ttgaactgtg gaaggagaag agctctggca 120

agtatgagaa gaaggtgtat gtgtctccca agcaccttga tgagaagggt gctgtctctcc 180
 acttggggcaa gcttgggtgcc aagctgacca agctcaccaa gtctcaggcc gactacatca 240
 gcgtgccgat cgagggtccc tacaagcctg cccat 275

<210> 1499
 <211> 306
 <212> nucleic acid
 <213> Zea mays

<400> 1499

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 gcggacttctg gcgcgcctcga gattgagctg gccgaggtcg aaatgcccg cctcatggcg 180
 tgccgcgcgcg agttcggccc gtccaagccc ttgcgcggcg ctaggatctc ggggtctctc 240
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 cgctgg 306

<210> 1500
 <211> 280
 <212> nucleic acid
 <213> Zea mays

<400> 1500

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 attgatatgc tcggccttga gacctaccct ggcgtcaagc gcatcaccat caagncccag 180
 actgaccgct ggggtgttccc cgagaccaac actggcatca ttgtccttgt tgaggggtgc 240
 tgatgaactt nggggtgcaa ttggccatcc caactttggc 280

<210> 1501
 <211> 293
 <212> nucleic acid
 <213> Zea mays

<400> 1501

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tgatgaacct tgggtgtgct aatntgccat cctaagcttg tcatgtcctg ctcatcact 120
aaccaggtca ttgcccaact tgaactgtgg aaggagaaga gctctggcaa gtatgagaag 180
aaggtgtatg tgctcccaaa gcaccttgat gagaaggttg ctgctctcca cttgggcaag 240
cttgggtgcca agctgaccaa gctcaccaag tctcaggccg actacatcag cgt 293

<210> 1502
<211> 307
<212> nucleic acid
<213> Zea mays

<400> 1502

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ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120
tctcgcaggc ggacttcggc cgctcgcaga ttgagctggc cgaggtcgaa atgcccggcc 180
tcatggcgtg ccgcgcgcag ttccggcccg ccaagccctt cgccggcgct aggatctcgg 240
ggctctctca catgaccatc cagaccgccc tctcatcga gacctcacc gcgctcggcg 300
ccgaggt 307

<210> 1503
<211> 232
<212> nucleic acid
<213> Zea mays

<400> 1503

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acaatgccat tgtctgcaac attggccact ttgacaatga aattgatatg ctgggccttg 120
agacctacc tggcgtcaag cgcacacca tcaagcccca gactgaccgc tgggtgttcc 180
ccgagaccaa cactggcatc attgtccttg ctgagggtcg cctgatgaac ct 232

<210> 1504
<211> 277
<212> nucleic acid
<213> Zea mays

<400> 1504

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ccgagttcgg ccggtccaag nccttcgccg gcgctaggat ctcggggtct ctccacatga 120
ccatccagac cgcggtcctc atcgagaccc tcaccgcgct cggcgccgag gtccgctggt 180
gtcctgcaa natcttctcc acgcaggacc acgtgccgc ngccatcgcg agcaantcgg 240
ccgngtntt cgcttaaang gggaaaccct cgngnat 277

<210> 1505
<211> 234
<212> nucleic acid
<213> Zea mays
<400> 1505

gttgtctctg aagotgacat ctctgtgacc accactggca acaaggatat catcatggtt 60
gaccacatga ggaanatgaa gaacaatgcc attgtctgca acattggcca ctttgacaat 120
gaaattgata tgctcggcct tgagacctac cctggcgctca agcgcatcac catcaagccc 180
cagactgacc gctgggtggt ccccgagacc aacactggca tcattgtcct tgct 234

<210> 1506
<211> 238
<212> nucleic acid
<213> Zea mays
<400> 1506

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agatgcagga gaccggcgcc ctctcttcc ctgccattaa cgtcaacgat tccgtcacca 120
agagcaagtt tgacaacctg tatggttgcc gccactcact ccctgatggt ctgatgaggg 180
ccaccgacgt tatgatcgcc ggtaagggtg ccgtggtctg cggatacggg gatgtttg 238

<210> 1507
<211> 281
<212> nucleic acid
<213> Zea mays
<400> 1507

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ccacgagggt gtcaaggccg aggaggatta cgagaagacc ggcaagatcc ccgaccggga 120
gtccaccgac aacgctgagt tcaagatcgt gtcaccatc atccgcgacg ggctcaaggc 180

tgaccccaag aagtaccgca agatgaagga gaggtctgtc ggcgtctctg aggagaccac 240
cacgggtgtc aagaggctct accagatgca ggagaccggc g 281

<210> 1508
<211> 235
<212> nucleic acid
<213> Zea mays

<400> 1508

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ccttgagacc taccctggcg tcaagcgcat caccatcaag cccagactg accgctgggt 120
gttccccgag accaactg gcatcattgt ccttgctgag ggtcgctga tgaaccttgg 180
gtgtgtact ggccatccta gctttgtcat gtctgtctca ttcactaacc aggc 235

<210> 1509
<211> 299
<212> nucleic acid
<213> Zea mays

<400> 1509

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cgtctggacg ggagtacaag gtcaaggatc tctcgaggc ggacttcggc cgctcgaga 120
ttgagctggc cgaggtegaa atgccgggc tcatggcgtg ccgcgccgag ttcggcccgt 180
ccaagccctt cgcggcgct aggatctcgg ggtctctcca catgaccatc cagaccgccg 240
tctcatcga gacctcacc gcgctcggcg ccgangtccg tgggtcctg caacatttc 299

<210> 1510
<211> 280
<212> nucleic acid
<213> Zea mays

<400> 1510

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gctcggcctt gagacctacc ctggcgtaaa gcgcatacc atcaagcccc agactgaccg 120
ctgggtgttc cccgagacca aactggcat cattgtcctt gctgagggtc gctgatgaa 180
ccttgggtgt gctactggcc atcctagctt tgtcatgtcc tgctcatcac taaccaggtc 240

atgcccaact tgaactgtgg aaggagaaga gctctggcaa 280

<210> 1511
 <211> 298
 <212> nucleic acid
 <213> Zea mays

<400> 1511

ctcgttccat ttcccatctt cccagatcca attcgcgagt tctccctcct ctgcggccat 60
 ggcgctctct gtggagaaga cctcgtctgg acgggagtag aaggtcaagg atctctcgca 120
 ggcggacttc ggccgcctcg agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180
 gtgccgcgcc gagttcggcc cgtccaagcc cttcgcgggc gctaggatct cggggtctct 240
 ccacatgacc atccagaccg cctcctctat cgagaccctc accgcgctcg gcgccgag 298

<210> 1512
 <211> 250
 <212> nucleic acid
 <213> Zea mays

<400> 1512

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 gaaattgata tgctcggcct tgagacctac cctggcgta agcgcatcac catcaagccc 120
 nagactgacc gctgggtggt ccccgagacc aacactggca tcattgtcct tgctgagggt 180
 cgctgatga acctgggtg tgctactggc catcctagct tggncatgtc cnnngctaann 240
 antaacnagg 250

<210> 1513
 <211> 291
 <212> nucleic acid
 <213> Zea mays

<400> 1513

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 ggggcgaggc gggcgccccc gacctcatcg tcgacgacgg cggcgacgcc acgctgctca 120
 tccacgaggg tgtcaaggcc gaggaggatt acgagaagac cgnccangatc cccgaccggg 180
 agtccaccga caacgctgag ttcaagatcg tgctcaccat catccgcgac gggctcaagg 240

ctgaccccaa gaagtaccgc aagatgaagg agaggcttgt cggcgtctct g 291

<210> 1514
<211> 300
<212> nucleic acid
<213> Zea mays

<400> 1514

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tggcgtctctc tgtggagaag acctcgtctg gacgggagta caaggtcaag gatctctcgc 120
aggcggactt cggcgcgcctc gagattgagc tggccgaggt cgaaatgcc ggccatcatgg 180
cgtgccgcgc cgagttcggc cegtccaagc ccttcgccgg cgctaggatc tcgggggtctc 240
tccacatgac catccagacc gccgtctctca tcgagaccct caccgcgctc ggccgccgagg 300

<210> 1515
<211> 237
<212> nucleic acid
<213> Zea mays

<400> 1515

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gacgttgtct ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg 120
gttgaccaca tgaggaagat gaagaacaat gccattgtct gcaacattgg ccactttgac 180
aatgaaattg atatgctcgg ccttgagacc taccctggcg tcaagcgcac caccatc 237

<210> 1516
<211> 245
<212> nucleic acid
<213> Zea mays

<400> 1516

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ctgaggggtcg cctgatgaac cttgggtgtg ctactggcca tcttagcttt gtcattgtcct 120
gtcattcac taaccaggtc attgccaac ttgaactgtg gaaggagaag agctctggca 180
agtatgagaa gaaggtgtat gtgctcccca agcaccttga tgagaagggt gctgctctcc 240
acttg 245

<210> 1517
 <211> 298
 <212> nucleic acid
 <213> Zea mays

 <400> 1517

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 atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg 120
 caggcggaact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
 gcgtgccgcy ccgagttcgg cccttccaag cccttcgccg gcgctaggat ctcggggtct 240
 ctccacatga ccattccagac cgcgtcctc atcgagacc tcaccgcgt cggcgccg 298

<210> 1518
 <211> 239
 <212> nucleic acid
 <213> Zea mays

 <400> 1518

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 gatcgtgctc accatcatcc ggcacgggct caaggctgac cccaagaagt accgcaagat 120
 gaaggagagg cttgtcggcg tctctgagga gaccaccacg ggtgtcaaga ggctctacca 180
 gatgcaggag accggcgccc tctcttccc tgccattaac gtcaacgatt ccgtcacca 239

<210> 1519
 <211> 278
 <212> nucleic acid
 <213> Zea mays

 <400> 1519

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 ctaaccaggt cattgcccac cttgaactgt ggaaggagaa gagctctggc aagtatgaga 120
 agaagggtgta tgtgtctccc aagcaccttg atgagaaggt tgctgtctc cacttgggca 180
 agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc agcgtgccga 240
 tcgagggtcc ctacaagcct gccactacc ggtactag 278

<210> 1520

<211> 272
 <212> nucleic acid
 <213> Zea mays

 <400> 1520

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 gaagatgaag aacaatgcca ttgtctgcaa cattggccac ttgacaatg aaattgatat 120
 gctcggccctt gagacctacc ctggcgtcaa gcgcatacc atcaagcccc agactgaccg 180
 ctgggtgttc cccgagacca aactggcat cattgtcttg ctgagggctg ctgatgaact 240
 tgggtgtgtat ggccatctag tttgtcatgt ct 272

<210> 1521
 <211> 283
 <212> nucleic acid
 <213> Zea mays

 <400> 1521

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 gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc ggacttcggc 120
 cgctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcatggcgtg ccgcgacgag 180
 ttcggcccggt ccaagccctt cgccggcgct aggatctcgg ggtctctcca catgaccatc 240
 cagaccgccc tctcatcga gacctcacc gcgctcggcg ncg 283

<210> 1522
 <211> 235
 <212> nucleic acid
 <213> Zea mays

 <400> 1522

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 cgagaccaac actggcatca ttgtccttgc tgagggtcgc ctgatgaacc ttgggtgtgc 120
 tactggccat cctagctttg tcatgtcctg ctattcact aaccagggtca ttgcccact 180
 tgaactgtgg aaggagaaga gctctggcaa gtatgagaag aagggtgtatg tgctc 235

<210> 1523
 <211> 313
 <212> nucleic acid

<213> Zea mays

<400> 1523

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gcgacgccac gctgctcacc cacganggtg tcaaggccga ggaggattac gagaagaccg 120
gcaaaatncc cgancccgga gatcancgga naaagcgngt ncaaaaagtg gnncaanaat 180
aatcggggaag gggtnaangn tnaccccnag aagtaccgca agatgaanga gaggcttgtn 240
ggcgtctctn aggagaccac cacgggtgtn aagaggctct accagatgca ggagaccggc 300
ggcctctctt tcc 313

<210> 1524

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1524

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ggcgctctct gtggagnaga cctcgtnnng acgggagtag aaggtcaagg atctctcgca 120
ggcggacttc ggccgcgcgc agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180
gtgccgcgcc gaggctcgcc cgtccaagcc ctccgccggc gctaggatct cgggggtctct 240
ccacatgacc atccagaccg ccgtctctcat cgagaccctc ancgcgctcg gcgcgcagg 299

<210> 1525

<211> 232

<212> nucleic acid

<213> Zea mays

<400> 1525

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caagatcgtag ctacccatca tccgcgacgg gctcaaggct gacccaaga agtaccgcaa 120
gatgaaggag aggcttgtag gcgtctctga ggagaccacc acgggtgtca agaggctcta 180
ccagatgcag gagaccggcg cctcctcttt cctgccatt aacgtcaacg at 232

<210> 1526

<211> 317

<212> nucleic acid

<213> Zea mays

<400> 1526

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 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120
 tctcgcaggc ggacttcggc cgcctcgaga ttgagctggc cgaggtcgaa atgcccggcc 180
 tcatggcgtg ccgcgcgcgag ttcggcccggt ccaagccttt cgcgcgcgct aggatcttcg 240
 gggctctctc acatgaccat ccagaccgcc gtctctcatg agaccctcac cgcgctggcg 300
 ccgaagtccg ctggtgt 317

<210> 1527

<211> 289

<212> nucleic acid

<213> Zea mays

<400> 1527

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 aggcggactt cggcgcgctc gagattgagc tggccgaggt cgaaatgcc gccctcatgg 180
 cgtgccgcgc cgagttcggc ccgtccaagc ctttcgccgg cgctaggatc tcgggggtctc 240
 tccacatgac catccagacc gccgtctctc tcgagaccct caccgcgct 289

<210> 1528

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1528

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 ggccatggcg ctctctgtgg agaagacctc gtctggacgg gagtacaagg tcaaggatct 120
 ctcgcaggcg gacttcggcc gcctcgagat tgagctggcc gaggtcgaaa tgcccgccct 180
 catggcgtgc ngcgacgagt tcggcccgct caagcccttc gccggcgcta ggatctcggg 240
 gtctctccac atgaccatcc agaccgccgt cctcatcgag accctcaccg cgctcggng 299

<210> 1529

<211> 245
 <212> nucleic acid
 <213> Zea mays

 <400> 1529

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 aaccagggtca ttgcccaact tgaactgtgg aaggagaaga gctctggcaa gtatgagaag 120
 aagggtgatg tgctcccaaa gcaacttgat gagaagggtg ctgctctcca cttgggcaag 180
 cttggtgcc aagctgaccaa gctcaccaag tctcaggccg actacatcag cgtgccgatc 240
 gaggg 245

<210> 1530
 <211> 287
 <212> nucleic acid
 <213> Zea mays

 <400> 1530

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 ggcgtctct gtggagaaga cctcgtctgg acgggagtag aaggtaagg atctctcgca 120
 ggcggacttc ggccgcctcg agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180
 gtgccgcgcc gagttcggcc cgtccaagcc cttcgccggc gctaggatct cgggggtctct 240
 ccacatgacc atccagaccg ccgtcctcat cgagaccctc accgcgc 287

<210> 1531
 <211> 283
 <212> nucleic acid
 <213> Zea mays

 <400> 1531

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 naggtttgtt cggcgtctct gaggagacca ccacgggtgt tcaagaggct ctaccagatg 120
 caggagaccg ggcacctct cttccctgcc attaacgtca acgattccgt caccaagagc 180
 aagtttgana acctgtatgg ttgccgcan tcgtcctgat ggtctgatga gggcactgac 240
 gttatgatcg ccggaaggtn gccgtggtct gcgaatacgt ntt 283

<210> 1532

<211> 301
 <212> nucleic acid
 <213> Zea mays

 <400> 1532

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 gcgtctctctg tggagaagac ctggtctgga cgggagtaca aggtcaagga tctctcgcag 120
 ggcgacttcg gccgcctcga gattgagctg gccgaggtcg aaatgcccg gctcatggcg 180
 tgccgcgncg agttcggccc gtccaagccc ttgcgcggcg ctaggatctc ggggtcttcc 240
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 t 301

<210> 1533
 <211> 268
 <212> nucleic acid
 <213> Zea mays

 <400> 1533

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 tgctcattca ctaaccaggt cattgcccaa cntgaactgt ggaaggagaa gagctctggc 120
 aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgctgctctc 180
 cacttgggca agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 240
 agcgtgccga tcgaggtcct acaagcct 268

<210> 1534
 <211> 286
 <212> nucleic acid
 <213> Zea mays

 <400> 1534

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 caggcggact tcggccgcct cgagattgag ctggccgang tcgaaatgcc cggcctcatg 180
 gcgtgccgcg ccgagttcgg cccgtccaan ccttcgccg gcgctaggat ctcggggtct 240
 ctccacatga ccatccagac cgccgtcctc atcgagacct tcaccg 286

<210> 1535
 <211> 233
 <212> nucleic acid
 <213> Zea mays

 <400> 1535

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 ccgacgttat gatcgccggt aaggttgccg tggctctgcg atacggtgat gttggcaagg 120
 gttgtgccgc tgcantcaag caggctggtg cccgtgtcat tgtgaccgag atcgacccca 180
 tctnngccct ccaggctctg ntggagggtc ttcaggctct tcccttggag gac 233

<210> 1536
 <211> 339
 <212> nucleic acid
 <213> Zea mays

 <400> 1536

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 gacctgtct ggacgggagt acaaggtcaa ggatctctcg caggcggact tcggccgcct 120
 cgagattgag ctggccgagg tcgaaatgcc cggcctcatg gcgtgccgcg ccgagttcgg 180
 ccgtccaag ccttcgccg gcgctaagat ctcggggtct ctccacatga ccatccagac 240
 cgcgctctc atcgagaacc tcaccgcntt ggccggaag tccgtggtgt cctgcaanat 300
 tttccangca gaccaggcgg cggcggcatg ggcggatgg 339

<210> 1537
 <211> 263
 <212> nucleic acid
 <213> Zea mays

 <400> 1537

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 gtcggcgtct ctgaggagac nccacggga nttcaagagg cactacncag atgcaggana 120
 ccggcgcct cctcttccct gccattaacg tcaacgattc cgtcaccaag agcaagtttg 180
 acaacctgta tggttgccgc cactcgctcc ctgatggtct gatgagggcc actgacgtta 240
 tgatcgccgg aaaggttgcc gtg 263

<210> 1538
 <211> 226
 <212> nucleic acid
 <213> Zea mays

 <400> 1538

 attgtgaccg agatogaccc catctgtgcc ctccaggctc tgatggaggg tottcaggtc 60
 cttcccttgg aggagtttgt ctctgaagct gacatcttcg tgaccaccac tggcaacaag 120
 gatatcatca tggttgacca catgaggaag atgaagaaca atgccattgt ctgcaacatt 180
 ggccactttg acaatgaaat tgatatnctc ggccttgata cctacc 226

<210> 1539
 <211> 302
 <212> nucleic acid
 <213> Zea mays

 <400> 1539

 ctcactcccg ttcentttcc ccatctccca gntccanttc gcgagttctc cctcctctgc 60
 cgccatggcg ntctctgtgg agaagacntc gtctggacgg gagtacaagg tcaaggatnt 120
 ctgcgaggcg ganttcggcc gctctgagat tganctggcc gaggtcgaaa tgcccggcct 180
 catggcggtgc cgcgcgaggt tgggcccgtc caancccttc gccggcncta ggatntcggg 240
 gtctctccac atgaccatcc agaccgcggt cctcatcgag accctcaccg cgtctggcnt 300
 ga 302

<210> 1540
 <211> 277
 <212> nucleic acid
 <213> Zea mays

 <400> 1540

 agctgacatc ttctgtgacca ccaactggcaa caaggatatc atcatgggttg accacatgag 60
 gaagatgaag aacaatgcca ttgtctgcaa cattggccac tttgacaatg aaattgatat 120
 gctcggcctt gagacctanc ctggcgtaaa ggcatcacca tcaagcccca gactgancgc 180
 tgggtgtttc cccgagacca aactggcat cattgtcctt gctgagggtc gctggatgna 240
 ncttgggtgt gctactgggc atcctagttt tgtcatg 277

<210> 1541
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 1541

 ccgttccatt tccccatctc ccagatccaa ttgcgcgagtt ctccctcctc tgccctcgact 60
 ggggcgagggc gggcgggccc gacctcatcg tcgacgacgg cggcgacgcc aacgtgtctc 120
 atccacgagg gtgtcaaggc cgaggaggat tacgagaaga ccggcaagat tccgacccgg 180
 agtccaccga caacgtgtgag ttcaagatcg tgctcaccat catccgcgac gggctcaagg 240
 ctgaccccaa gaagtaccgc aagatgaagg agag 274

<210> 1542
 <211> 243
 <212> nucleic acid
 <213> Zea mays

 <400> 1542

 cggaacgctg ggangctgac atcttcgtga ccaccactgg caacaaggat atcatcatgg 60
 ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc cactttgaca 120
 atgaaattga tatgtctggc cttgagacct accctggcgt caagcgcac accatcaagc 180
 cccagactga ccgtgggtgt tccccgagac caacactggc atcatgtctt gtgaaggctg 240
 ctg 243

<210> 1543
 <211> 284
 <212> nucleic acid
 <213> Zea mays

 <400> 1543

 cctcactccc gttccatttc cccatctccc agatccaatt cgcgagttct cctcctctg 60
 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120
 tctgcaggc ggacttcggc cgctctgaga ttgagctggc cgaggtcgaa atgcccggcc 180
 tcatggcgtg ccgcgcgag ttccggcccgt ccaagccctt cgccggcgct aggatctcgg 240
 ggtctctcca catgaccatc cagaccgccg tctcatcga gacc 284

<210> 1544
 <211> 261
 <212> nucleic acid
 <213> Zea mays

 <400> 1544

 ctcccagatc caattcgcca gttctccctc ctctgccgcc atggcgctct ctgtggagaa 60
 gacctcgtct ggacgggagt acaagggtcaa ggatctctcg caggcggact tcggccgcct 120
 cgagattgag ctggccgagg tcgaaatgcc cggcctcatg gcgtgccgcg ccgagttcgg 180
 cccgtccaag cccttcgccg gcgctaggat ctcggggtct ctccacatga ncatccagac 240
 cgccgtcctc atcgagaccc t 261

<210> 1545
 <211> 280
 <212> nucleic acid
 <213> Zea mays

 <400> 1545

 ctctcgttcc atttccccat ctcccagatc caattcgcca gttctccctc ctctgccgcc 60
 atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg 120
 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
 gcgtgccgcg ccgagttcgg cccgtccaag cccttcgccg gcgctaggat ctcggggtct 240
 ctccacatga ccatccagac cgccgtcctc atcgagaccc 280

<210> 1546
 <211> 288
 <212> nucleic acid
 <213> Zea mays

 <400> 1546

 ctccctcact cccgttccat ttccccatct cccagatcca attcgcgagt tctccctcct 60
 ctgccgccat gcgctctctg tggagaagac ctcgctctga cgggagtaca aggtcaagga 120
 tctctcgcag gcggaattcg gccgcctcga gattgagctg gccgaggtcg aaatgcccg 180
 cctcatggcg tgccgcgccg agttcgcccc gtccaagccc ttgcggcgcg ctaggatctc 240
 ggggtctctc cacatgacca tccagaccgc cgtcctcatc gagaccct 288

<210> 1547
 <211> 260
 <212> nucleic acid
 <213> Zea mays

 <400> 1547

 atctcccaga tccaattcgc gagttctccc tctctgcgg ccattggcgt ctctgtggag 60
 aagacctcgt ctggacggga gtacaaggtc aaggatctct cgcaggcgga cttcggccgc 120
 ctcgagattg agctggccga ggtcgaaatg cccggcctca tggcgtgccg cgccgagttc 180
 ggcccgtcca agcccttcgc cggcgctagg atctcggggt ctctccacat gaccatccag 240
 accgcngtcc tcatcgagac 260

<210> 1548
 <211> 212
 <212> nucleic acid
 <213> Zea mays

 <400> 1548

 tactggtggt gcaaccgagcg ctgcctcgac tggggcgagg cgggcggcca cgacctcacc 60
 gncgacgacg gcggcgacgc caccgtgctc atccacgagg gtgtcaaggc cgaggaggat 120
 tacgagaaga ccggcaagat ccccgacccg gagtcancgg acaacgctga gttcaagatc 180
 gtgctcacca tcatccgcga cgggctcaag gt 212

<210> 1549
 <211> 277
 <212> nucleic acid
 <213> Zea mays

 <400> 1549

 cggacgntgg tagtggttgt gccgctgcac tncaagcagg ctggtgcccg tgtcattgng 60
 nccgagatcg accccatctg cncctccag gctctgatgg agggctctca ggtccttccc 120
 ttggaggacg ttgtctcgga agctgacatc ttcgtgacca ccaactggcaa caaggatatc 180
 atcatggttg accacatgag gaagatgaag aacaatgcc a ttgtctgcaa cattggccat 240
 ttgacaatga attgatatgc tcggccttga gacctac 277

<210> 1550

<211> 277
 <212> nucleic acid
 <213> Zea mays

 <400> 1550

 ggtgcaccga ggcgtgcctc gactggggcg angcgngcgg cnccgacctc atcgtcgacg 60
 acggcngcga cgccacgctg ctcanccacg anggtgtcaa ggccgntggg gattacgagn 120
 agaccggcna gatccccgac ccgngtnca ccgacaacgc tgagttcaag atcggtctca 180
 ccatcntccg ngacgggctn aacgtgacc ccaagaagta ccgcnaantg aangagangt 240
 tgtaggcgtc totgangaga ncacnacggg tgtnaag 277

<210> 1551
 <211> 291
 <212> nucleic acid
 <213> Zea mays

 <400> 1551

 ctttgtcatg tctgtctcat tcactaacca ggtcattgcc naacttgaac tgtggaagga 60
 gaagagctct ggcaagtatg agaagaaggt gtatgtgctc cccaagcacc ttgatgagaa 120
 gggttgetgt ctccacttgg gcaagcttgg tgccaagctg accaagctca ccaagtctca 180
 ggccgactac atcagcgtgc cgatcgaggg tccttacaag cctgcccact accggtacta 240
 ggcacacggc ttgcagctca ctcgggccgt tgtgtgctat gaagttcgct a 291

<210> 1552
 <211> 274
 <212> nucleic acid
 <213> Zea mays

 <400> 1552

 ctcccgttcc atttccccat ctcccagatc caattcgca gttctccctc ctctgccgcc 60
 atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg 120
 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
 gcgtgccgcg ccgagttcgg ccggtccaag cccttcgccg gcgctaggat ctgggggtct 240
 ctccacatga ccatccagac cgccgtcctc atcg 274

<210> 1553

<211> 318
 <212> nucleic acid
 <213> Zea mays

 <400> 1553

 tagctttgtc atgtcctgct cattcactaa ccaggtcatt gcccaacttg aactgtggaa 60
 ggagaagagc tctggcaagt atgagaagaa ggtgtatgtg ctccccaagc accttgatga 120
 gaaggttgct gctctccact tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc 180
 tcaggccgac tacatcagcg tgccgatcga gggtcctaa caagcctgcc cactaccggt 240
 actaggcagc cagcacacgg cttgcagctc actcggggcg tgtgtgcaan nanttcogana 300
 cngggcctgn aatnattt 318

<210> 1554
 <211> 222
 <212> nucleic acid
 <213> Zea mays

 <400> 1554

 gcctgatgaa ccttgggtgt gctactggcc atcctagctt tgtcatgtcc tgotcattca 60
 ctaaccaggt cattgccc aa cttgaactgt ggaaggagaa gagctctggc aagtatgaga 120
 agaaggtgta tgtgctcccc aagcancttg atgagaaggt tgctgctctc cacttgggca 180
 agcttgggtgc caagctgacc aagttcacca agtctcaggc cg 222

<210> 1555
 <211> 259
 <212> nucleic acid
 <213> Zea mays

 <400> 1555

 catttcccca tctccagat ccaattcgcg agttctccct cctctgcggc catggcgctc 60
 tctgtggaga agacctcgtc tggacgggag tacaaggta aggatctctc gcaggcggac 120
 ttcgcccgcc tcgagattga gctggccgag gtcgaaatgc ccggcctcat ggcgtgccgc 180
 gccgagttcg gcccgccaa gcccttcgcc ggcgctagga tctcggggtc tctccacatg 240
 accatccaga ccgccgtcc 259

<210> 1556

<211> 260
 <212> nucleic acid
 <213> Zea mays

 <400> 1556

 cgttccatTT ccccatctcc cagatccaat tcgcgagttc tcctctctct gcggccatgg 60
 cgctctctgt ggagaagacc tcgtctggac gggagtacaa ggtcaaggat ctctcgcagg 120
 cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggc ctcatggcgt 180
 gccgcgccga gttcggcccc tccaagccct tcgccggcgc taggatctcg gggctctctcc 240
 acatgaccat ccagaccgcc 260

<210> 1557
 <211> 271
 <212> nucleic acid
 <213> Zea mays

 <400> 1557

 cctcactccc gtTccatttc cccatctccc agatccaatt cgcgagttct ccctcctctg 60
 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120
 tctcgcaggc ggacttcggc ccgcctcgaga ttgagctggc cgaggtcgaa atgcccggcc 180
 tcatggcgtg ccgcgccgag ttccggcccc ccaagccctt cgcgggcgct aggatctcgg 240
 ggtctctcca catgaccatc cagaccgccc c 271

<210> 1558
 <211> 223
 <212> nucleic acid
 <213> Zea mays

 <400> 1558

 ttctcatgt ggtcaaccat gatgaagatg aagaacaatg ccattgtctg caacattggc 60
 cactttgaca atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcac 120
 accatcaagc ccagactga ccgctgggtg ttccccgaga ccaacactgg catcattgtc 180
 cttgctgagg gtgcctgat gaacttgggt gtgctactgg cca 223

<210> 1559
 <211> 293
 <212> nucleic acid

<213> Zea mays

<400> 1559

cccgttccat ttccccatct cccagatcca attcgcgagt tctccctcct ctgccgccat 60
ggcgctgctc tgtggagaag acctcgctctg gacgggagta caangtcaag gatctctcgc 120
aggcggactt cggcgcgcctc gagattgagc tggccgaggt cgaaatgccc ggccatcatgg 180
cgtgccgcgc cgagttcggc cgtccaagc cttcgccggc cgttaggata tcggggctctc 240
tccacatgac catccanacc gcngtctca tcgagacctc accgnttgg nnc 293

<210> 1560

<211> 244

<212> nucleic acid

<213> Zea mays

<400> 1560

cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60
cctcgctctgg acgggagtag aaggtcaagg atctctcgca ggccgacttc ggccgcctcg 120
agattgagct ggccgaggtc gaaatgcccc gcctcatggc gtgccgcgcc gagttcgggc 180
cgtccaagcc cttcgccggc gctaggatct cggggctctct ccacatgacc atccagaccg 240
cogt 244

<210> 1561

<211> 358

<212> nucleic acid

<213> Zea mays

<400> 1561

gaagngcaag tntgacnacc tgtatggttg ccgnactcg ctcccngatg gncatgatgan 60
ggcnactgac gttatgatcg nccgaaaagc tgccgtngtc ttgcgaatan gtnntgtggg 120
caaggtggtg ttnttgccct caagaaggcn gtgcccgtgt canttgaccg agatcgaccc 180
catctgtgcc tccaggctct gatggagggt cttcaggctt tcccttgag gagttgtct 240
ctgangtgac atcttcgtga ccaccactgg caaccaagga tatcncatgg ttgancacat 300
gagaagatga agaacaatgc cattgtctgc aacattggca cttgacaaga antgtatc 358

<210> 1562

<211> 218
 <212> nucleic acid
 <213> Zea mays

<400> 1562

gtcatgtcct gctcattcac taaccaggtc attgcctaac ttgaactgtg gaaggagaag 60
 agctctggca agtatgagaa gaagggtgtat gtgctcccca agcaccttga tgagaagggt 120
 gctgctctcc acttggggcaa gcttggtgcc aagctgacca agctcaccaa gtctcaggcc 180
 gactacatca gcgtgccgat cgagggtccc tacaagcc 218

<210> 1563
 <211> 269
 <212> nucleic acid
 <213> Zea mays

<400> 1563

cctccctcac tctcgttcca tttcccatc tcccagatcc aattcgcgag ttctccctcc 60
 tctgcggcca tggcgtctc tgtggagaag acctcgtctg gacgggagta caaggtaag 120
 gatctctcgc aggcggactt cggccgcctc gagattgagc tggccgaggt cgaaatgcc 180
 ggccatcatg cgtgccgcgc cgagttcggc ccgtccaagc ccttcgccgg cgctaggatc 240
 tcggggtctc tccacatgac catccagac 269

<210> 1564
 <211> 260
 <212> nucleic acid
 <213> Zea mays

<400> 1564

ctctcgttcc atttcccat ctcccagatc caattcgcga gttctccctc ctctgcggcc 60
 atggcgtctc ctgtggagaa gacctcgtc ggacgggagt acaaggtaaa ggatctctcg 120
 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
 gcgtgccgcg ccgagttcgg ccgtccaag cccttcgccg gcgctaggat ctcggggtct 240
 ctccacatga ccatccagac 260

<210> 1565
 <211> 273
 <212> nucleic acid

<213> Zea mays

<400> 1565

cgactggggc gaggggggcg gccccgacct catcgtcgac gacggacggc gacgccacgc 60
tgctncattc cagcaggggtg tcaaggccga ggaggactac gagaagaccg gcaagatccc 120
cgatccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca tccgcgacgg 180
gctcaaggct gacccaaga agtaccgcaa gatgaaggag aggttgtcgg cntctctgag 240
gagaccacca cgggtgtcaa gaggctctac nat 273

<210> 1566

<211> 282

<212> nucleic acid

<213> Zea mays

<400> 1566

cattcactaa ccaggtcatt gcccaacttg aactgtggaa ggagaagagc tctggcaagt 60
atgagaagaa ggtgtatgtg ctccccaaagc accttgatga gaaggttgct gctctccact 120
tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc tcaggccgac tacatcagcg 180
tgccgatcga ggggtccctac aagcctgccc actaccggta ctaggcagcc agcacacggc 240
ttgcagctca ctggggccgt tgtgtgctat gaagttcgct ac 282

<210> 1567

<211> 235

<212> nucleic acid

<213> Zea mays

<400> 1567

ctcgcaggcg gacntcggcc gcctcgagat tgagcnggcc gaggtcgaaa tgcccggcct 60
catggcgtgg ccgcgccgag ttcggcccga ncaaagcnc tgcgccggcg taggatcncg 120
gggtctcncc acatgaccat ccagaccgcc gtctcctcgc agaccntcac ngcgtcggc 180
gccgaggtcc gtggtgcnc ngcaacatct tctccagcag gancacgccg ccgcc 235

<210> 1568

<211> 239

<212> nucleic acid

<213> Zea mays

<400> 1568

cttcggccgc ctcgagattg agctggccga ggtcgaaatg cccggcctca tggcgtgccg 60
 cgccgagttc ggcccgctcca agcccttcgc cggcgctagg atctcggggg ctctccacat 120
 gaccatccag accgccgtcc tcatcgagac cctcaccgcg ctcggcgcgc aggtcngctg 180
 gtgtccgca acatctttca acgcaggaca aggcngcggc ggcatcggcg gattcgggtg 239

<210> 1569

<211> 250

<212> nucleic acid

<213> Zea mays

<400> 1569

gttcacatttc cccatctccc agatccaatt cgcgagttct cctcctctg cggccatggc 60
 gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc 120
 ggacttcggc cgctcgcaga ttgagctggc cgaggtcgaa atgcccggcc tcattggcgtg 180
 ccgcgcgcgag ttcgggccgt ccaagccctt cgcgcggcgt aggatctcgg ggtctctcca 240
 catgaccatc 250

<210> 1570

<211> 264

<212> nucleic acid

<213> Zea mays

<400> 1570

ctccctcaact cccgttccat ttccccatct cccagatcca attcgcgagt tctccctcct 60
 ctgccgccat ggcgctctct gtggagaaga cctcgtctgg acgggagtac aagggtcaagg 120
 atctctcgca ggcggacttc ggccgcctcg agattgagct ggccgaggtc gaaatgcccg 180
 gctcatggc gtgcgcgcgc gagttcggcc cgtccaagcc cttcgccggc gctaggatct 240
 cgggggtctct ccacatgacc atcc 264

<210> 1571

<211> 274

<212> nucleic acid

<213> Zea mays

<400> 1571

tttactaac caggtcattg cccaacttga actgtggaag gagaagagct ctggcaagta 60
 tgagaagaag gtgtatgtgc tccccaaagca ccttgatgag aaggttgctg ctctccactt 120
 gggcaagctt ggtgccaagc tgaccaagct caccaagtct caggccgact acatcagcgt 180
 gccgatcgag ggtccctaca agcctgcccc ctaccggtac taggcnacgg cttgcagctt 240
 cactcgggcc gttgtgtgct atgaagttcg ctac 274

<210> 1572
 <211> 289
 <212> nucleic acid
 <213> Zea mays

<400> 1572

tttccccatc tcccagatcc aattcgcgag ttctccctcc tctgccgcca tggcgtcttc 60
 tgtggagaag acctcgtctg gacgggagta cataggtcaa ggatctctcg caangcggga 120
 cttcggccgc ctgagattg agctggccga ggtcgaaatg cccggnetca tggcgtgccg 180
 cgccgagttc ggcccgctca ancccttcgc cggcgctaag atctcggggg ctctccacat 240
 gaccatccag ancgncgtcc tcatcgagac cctcancng ctngggggg 289

<210> 1573
 <211> 276
 <212> nucleic acid
 <213> Zea mays

<400> 1573

atcaagcccc agactgaccg ctgggtgttg atggtgatgc gcttgacgcc agggtaggtc 60
 tcaaggccga gcatatcaat ttcatgtca aagtggccaa tgttgacagc aatggcattg 120
 ttcttcatct tcctcatgtg gtcaaccatg atgatatcct tgttgccagt ggtggtcacg 180
 aagatgtcag cttccgaggt caacggctcc aagctgtcgc ctgangagct cgtggtgctg 240
 caggggtgcc cgtgcccgcc gtctaagctc cgcccc 276

<210> 1574
 <211> 310
 <212> nucleic acid
 <213> Zea mays

<400> 1574

ctcntcactc tegtccatt tccccatctc ccagatccaa ttcgaggtt ctccctctc 60
 tgcggccatg gogctctctg tggagaagac ctggtctgga cgggagtaca angtaagga 120
 tctctgcag ggggaattcg gccgcctcga gattgagctg gccgaggtcg aaatgcccg 180
 cctcatggcg tgcgcgcgcg agttcggccc gtccaagcnc ttcgccggcg taggattcgg 240
 ggtctctcca catgaccatc cagaccgcg tctcatcgag acctcacgcg tcggcgccga 300
 ggtcgtggtg 310

<210> 1575
 <211> 241
 <212> nucleic acid
 <213> Zea mays

<400> 1575
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 gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgaggc ggacttcggc 120
 cgcctcgaga ttgagctggc cgaggtcgaa atgcccgcc tcattggcgtg ccgcgccgag 180
 ttcggcccggt ccaagccctt cgcgggcgct angatctcgg ggtctctcca catgacatcc 240
 a 241

<210> 1576
 <211> 299
 <212> nucleic acid
 <213> Zea mays

<400> 1576
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 gaagagctct ggcaagtatg agaagaaggt gtatgtgctc cccaagcacc ttgatgagaa 120
 ggttgtctgt ctccacttgg gcaagcttgg tgccaagctg accaagctca ccaagtctca 180
 ggccgactac atnccgcgtg ccgatcgagg gtccctacaa gcctgcccac ttaccggtat 240
 aggcacacgg cttgcagctc actcggggcg ttgtgtgcta tgaattcgct acatggctg 299

<210> 1577
 <211> 272
 <212> nucleic acid
 <213> Zea mays

<400> 1577

ccgacgctgg ncggnacgcg tccggttcca tttcccatc tcccagatcc aattcgcgaa 60
gntccctcc tctgcgcca nggcgtctc tgtggagaag acctcgtctg gacgggagta 120
caaggtoaag gatctctcgc aggcggactt cggccgcctc gagattgagc tggccnaggt 180
cgaaatgcc ggctcatgg cgtgccgcgc cgagttcggc ccgtccaagc ccttcgccgg 240
cgctaggatc tcggggtctc tccacatgac ca 272

<210> 1578

<211> 179

<212> nucleic acid

<213> Zea mays

<400> 1578

ggagaccgga gccctctct tccctgccat taacgtcaac gattccgtca ccaagagcaa 60
gtttgacaac ctgtatggtt gccgccactc actccctgat ggtctgatga gggccaccga 120
cgttatgac gccggtaagg ttgccgtggt ctgccgatac ggtgatgttg gcaagggtt 179

<210> 1579

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1579

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tggagaagac ctcgtctgga cgggagtaca aggtcaagga tctctcgag gcggacttcg 120
gccgcctcga gattgagctg gccgaggtcg aaatgcccg cctcatggcg tgccgcgcgc 180
agttcgccc gtccaanccc ttngccggcg taggatctcg ggtcctcca catgaccatc 240
cagaccgng tentatcgag acctnacnng gttggngg 278

<210> 1580

<211> 243

<212> nucleic acid

<213> Zea mays

<400> 1580

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gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc ggacttcggc 120
cgctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcatggcgtg ccgcgccgag 180
ttcggcccgt ccaagccttt cgccggcgta ggatctcggg gtctctccac atgaccatcc 240
aga 243

<210> 1581
<211> 247
<212> nucleic acid
<213> Zea mays
<400> 1581

ctcgttccat tccccatct cccagatcca attcgcgagt tctccctcct ctgcggccat 60
ggcgtctctt gtggagaaga cctcgtctgg acgggagtac aagggtcaagg atctctcgca 120
ggcggacttc ggcgcctcg agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180
gtgcgcgcgc gagttcggcc cgtccaagcc cttcgcggcg gctaggatct cggggtctct 240
ccacatg 247

<210> 1582
<211> 246
<212> nucleic acid
<213> Zea mays
<400> 1582

cgttccatt tccccatct cccagatcca ttcgcgagtt ctccctcctc tgccgccatg 60
gogctctctg tggagaagac ctcgtctgga cgggagtaca aggtcaagga tctctcgcag 120
goggacttcg gccgcctcga gattgagctg gccgaggtcg aaatgcccg cctcatggcg 180
tgccgcgcgc agttcggccc gtccaagccc ttcgcggcg ctaggatctc ggggtctctc 240
cacatg 246

<210> 1583
<211> 276
<212> nucleic acid
<213> Zea mays
<400> 1583

cctccctcac tctcgttcca tttccccatc tcccagatcc aattcgcgag ttctccctcc 60

tctgcggccca tggcgctctc tgtggagaag acctcgctctg gacgggagta caaggtcaag 120
gatctctcgc agggcgactt cggccgcctc gagantgagc cggccgaggg cgaaaanccc 180
ggcctcaagg cgtgccgcgc cgagtnggcc cgnccaagcc cttcgccggc gctaggatct 240
cggggtctct ccacatgacc atccagaccg ccgtcc 276

<210> 1584
<211> 178
<212> nucleic acid
<213> Zea mays

<400> 1584

attgtotgca acattggcca ctttgacaat gaaattgata tgctcggcct tgagacctac 60
cctggcgtea agcgcatcac catcaagccc cagactgacc gctgggtgtt ccccagagacc 120
aacaactggca tcattgtect tgctgagggg cgctganga nccttgtgtg tactnntg 178

<210> 1585
<211> 175
<212> nucleic acid
<213> Zea mays

<400> 1585

cgacggcggc gacgccacgc tgctcatcca cgaggggtgtc aaggccgaga aggagtacga 60
gangaccggc aagatccccg acccgaggtc caccgacaac gctgagttca agatcgtgct 120
caccatcatc cgcgacgggc tcaaggctga cccaagaag taccgcaaga tgaag 175

<210> 1586
<211> 235
<212> nucleic acid
<213> Zea mays

<400> 1586

tccccatctc ccagatccaa ttcgagagtt ctccntcctc tgccgccatg gcgctctctg 60
tggaagaagac ctcgtctgga cgggagtaca aggtcaagga tctctcgag gcggacttcg 120
gccgcctcga gantgagctg gccgaggtcg aaatgcccg cctcatggcg tgccgcgccg 180
agttcgcccc gtccaagccc ttcgccggcg ctaggatctc ggggtctctc cacat 235

<210> 1587

<211> 180
 <212> nucleic acid
 <213> Zea mays

<400> 1587

ancaagtttg acaacctgta tggttgccgc nactcgtcc ctgatggtct gatgagggcc 60
 actgacgtta tgatcgccgg aaaggttgcc gtggtctgcg gatacgggtga tgtcgnnang 120
 ggttggtgng ctgcnctcaa gcaggctggt gcccggtgna tcgtgancgn gatcgacccc 180

<210> 1588
 <211> 236
 <212> nucleic acid
 <213> Zea mays

<400> 1588

cccgttocat ttcccatct cccagatcca attcgcgagt tctccctcct ctgccgccat 60
 ggcgctctct gtggagaaga cctcgtctgg acgggagtag aaggtcaagg atctctcgca 120
 ggcggacttc ggccgcctcg agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180
 gtgccgcgcc gaggtcggcc cgtccaagcc cttcgccggc gctaggatct cgggggt 236

<210> 1589
 <211> 272
 <212> nucleic acid
 <213> Zea mays

<400> 1589

ccttgggagg acgttgtctc ggaagctgac atcttcgtga ccaccactgg caacaaggat 60
 atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 120
 cacattgaca aagaaatgnt angcncgggc ccgnagaccn aacccggcgg caanngnanc 180
 aacaacnagg cccagacgga ncgcccgggg gnccccgaga ccaacacggn aacaagtcct 240
 gncgaggggc gcccgangaa ccatngggga gc 272

<210> 1590
 <211> 260
 <212> nucleic acid
 <213> Zea mays

<400> 1590

ccctcactcc cgttccattt ccccatctcc cagatccaat tcgcgagttc tccctcctct 60
 gccgccatgg cgtctctgtt ggagaagacc tcgtctggac gggagtacaa ggtcaaggat 120
 ctctgcagg cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggc 180
 ctcatggcgt gccgcgcga gttcggcccg tccaagcent tcgccggcgc taggatctcg 240
 ggggtctctc cacatganca 260

<210> 1591
 <211> 245
 <212> nucleic acid
 <213> Zea mays

<400> 1591

cctcactccc gttccttttc cccntctccc agntccantt cgcgagttct cctcctctg 60
 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggntc 120
 tctgcaggc ggaattcggc ccgctcgaga ttgagctggc cgaggtcga atgcccggcc 180
 tcatggcgtg ccgcgcgcga ttcggcccg ccaagccctt cgcggcgct aggatctcg 240
 ggtct 245

<210> 1592
 <211> 209
 <212> nucleic acid
 <213> Zea mays

<400> 1592

cccagatcca attcgcgagt nctccctcct ctgccgccat ggcgctctct gtggagaaga 60
 cctcgtctgg acgggagtac aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120
 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180
 cgtccaagcc cttcgcgggc gctaggatt 209

<210> 1593
 <211> 221
 <212> nucleic acid
 <213> Zea mays

<400> 1593

cccatctccc agatccaatt cgcgagttct cctcctctg ccgccatggc gctctctgtg 60

gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc ggacttcggc 120
 cgccctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcatggcgtg ccgcgccgag 180
 ttcggccccgt cnaagccctt cgcnggcggt aaggnnttgg g 221

<210> 1594
 <211> 226
 <212> nucleic acid
 <213> Zea mays

<400> 1594

ctctcgttcc atttcccat ctcccagatc caattcgca gttctccctc ctctcgggcc 60
 atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg 120
 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
 gcgtgcgcg ccgagttcgg ccggtccaag cccttcgccg gcgcta 226

<210> 1595
 <211> 149
 <212> nucleic acid
 <213> Zea mays

<400> 1595

ccctcgagga gtactggtgg tgcaccgagc gctgcctcga ctggggcgag gcggggcgcc 60
 ccgacctcat cgtcgacgac ggcggcgacg ccacgctgct catccacgag ggtgtcaagg 120
 ccgaggagga ttacgagaag accggcaag 149

<210> 1596
 <211> 301
 <212> nucleic acid
 <213> Zea mays

<400> 1596

cccgtncat ntncccatn ncccagatcc aatttcgca gtacnaccnc ctnagnccg 60
 catgggggct ctctgtggag aagacatcgt ctggacggga gtacaagggtc aaggatcnct 120
 cgcaggcgga cntcgccgc ntcgagattg agcnggccga ggtcgaaatn acccggnctc 180
 atggcgtnn gcgcgagtt cggcccgtcc aagcncttcg ccggcgctan gattcgggggt 240
 ctctaccaca tgaccatcca gacggcgctc tcatcgagac atcacagcgt cggngccgag 300

t

301

<210> 1597
 <211> 174
 <212> nucleic acid
 <213> Zea mays
 <400> 1597

ccacgctgct catccacgag ggtgtcaagg ccgaggagga gtacgagaag accggnaaga 60
 tccccgaccc ngngtccacc gataacnctg nnttcaagat cgtggctcac catcatccgc 120
 gacgggctca aggctgacnc caagaagtac cgcaagatga aggagnggct ngtt 174

<210> 1598
 <211> 228
 <212> nucleic acid
 <213> Zea mays
 <400> 1598

ccctcactcc cgttccattt ccccatctcc cagatccaat tcgcgagttc tccctcctct 60
 gcgcgccatgg cgctctctgt ggagaagacc tcgtctggac gggagtacaa ggtcaaggat 120
 ctctcgcagg cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggn 180
 ctcatggcgt gccgcgccga gttcggcccg tccaagccct tcgccggg 228

<210> 1599
 <211> 227
 <212> nucleic acid
 <213> Zea mays
 <400> 1599

cccagatcca attcgcgagt tctccctcct ctgcggccat ggcgctctct gtggagaaga 60
 cctcgtctgg acgggggtac aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120
 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180
 cgtccaagcc cttcgccggc gtaggatctc ggggtcctca natgaca 227

<210> 1600
 <211> 236
 <212> nucleic acid
 <213> Zea mays

<400> 1600

cggttccatt tccccatctc ccagatccaa ttcgcgagtt ctccctcctc tgccgccatg 60
gcgctctctg tggagaagac ctcgctctgga cgggagtaca aggtcaagga tctctcgcag 120
gcggaacttcg gccgcctcga gattgagctg gccgaggtcg aaatgcccgg cctcatggcg 180
tgccgcgcgcg agttcggccc gtccaagcct tcgcccggcg taggatctcg gggttc 236

<210> 1601

<211> 209

<212> nucleic acid

<213> Zea mays

<400> 1601

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atggcgctct ctgtggagaa gacctgctt ggacgggagt acaaggtaaa ggatctctcg 120
caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180
gcgtgcgcgcg ccgagttcgg cccgtccaa 209

<210> 1602

<211> 426

<212> nucleic acid

<213> Zea mays

<400> 1602

cctcaactcc gttccatttc cccanctccc agatccaatt cgcgagttct cccnctctg 60
ccgccanggn gctctctgtg gagaagacct cntctggacg ggagtacaag gtcaaggatn 120
nctcgcaggn ggacttcggt cggcncgaga ttganctggc ngagncgaa atgcccggcc 180
ncatggcntg ccgngccgan ttcggcccgt ctangccctt cgnccggcgt aggnnacng 240
ngtctacca catgaccatc naagactgtn ttcctcancg agaactnang gtacgacggt 300
accnaagtnc ccngatatcc gngnnnact tcccaacant tgaggaaact acantgcatt 360
tcnanngacc gnaccanaat tgncanacta aaggtaaatt aanncntagn ctncaanggg 420
naantg 426

<210> 1603

<211> 232

<212> nucleic acid

<213> Zea mays

<400> 1603

ccccatctcc cagatccaat tcgcganttcc tccctcctct gcggccatgg cgctctctgt 60
ggagaagacc tcgtctggac gggagtagaa ggtcaangat ctctcgcagg cggacttcgg 120
ccgntcnat attgagctgg ccgaggtaga aatgcncggc ctcatagcgt gccgcgccga 180
tttcggncog tcaagccct tcgcgggng ctaggatctc ggggtctctc ca 232

<210> 1604

<211> 218

<212> nucleic acid

<213> Zea mays

<400> 1604

ctccctcaen cccgttccan ttccccancc cccagancca attcgcgagt tctccctcct 60
ctgcgcgcac ggcgtctctc gtggagaaga cctcgtctgg acgggagtag aaggtagagg 120
atcnctcgca gggggacntc ggccgcctcg agantgagcn ggccgaggtc gaaatgcccg 180
gcctcatggc gtgcgcgcgc gagttcggcc cgtccaag 218

<210> 1605

<211> 134

<212> nucleic acid

<213> Zea mays

<400> 1605

gccgccactc actccctgat ggtctgatga gggccaccga cgttatgatc gccggtagg 60
ttgccgtggc ctgcggatag ggtgatgttg gcaagggttg tgccgctgca ctcaagcagg 120
ctggtgcccg tgtc 134

<210> 1606

<211> 152

<212> nucleic acid

<213> Zea mays

<400> 1606

cgccgtctc atcgagaccc tcaccgcgct cggcgccgag gtccgctggc gtcctgcaa 60
catcttctcc acgcaggacc acgccgccgc cgcnatcgcg cgcgantcgg ccgccgtgtt 120

cgctgnaaa gggggagacc cttgaggagt ag

152

<210> 1607
<211> 287
<212> nucleic acid
<213> Zea mays

<400> 1607

gtccaagtat gagaagaagg tgtatgtgct cccaagcac cttgatgaga aggntgctgc 60
nnctcccact tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc tcaggccgac 120
tacatcagcg tgccgatcga gggtccttac aagcctgccc actaccggta ctaggcagcc 180
agcacacggc ttgcagctca cttcggggccg ttgtgtgcta tnaagtnenc ncncaactgnc 240
ctgtcagttc atcttttgca tgcatatgca ntatcatata cgcacg 287

<210> 1608
<211> 123
<212> nucleic acid
<213> Zea mays

<400> 1608

tgtctctgaa gctgacatct tcgtgaccac cactggcaac aaggatatca tcatgggttga 60
ccacatgagg aagatgaaga acaatgccat tgtctgcaac attggccact ttgacaatga 120
ant 123

<210> 1609
<211> 348
<212> nucleic acid
<213> Zea mays

<400> 1609

atcgccggaa aggttgccgn ggtctgcgga tacggtgatg tnnncaaggg ttgtgctgct 60
gccctcaagc aggttggtgc ccgtgtcatt gtgaccgaga tcgaccccat ctgtgcnctc 120
caggctctga tggagggctt caggtcttcc cttggaggac gtgtctctga agcnnacatc 180
tcgtgacaac attggcanca agtatcatca tggtgaccac atgaggaaga gaagacccat 240
gccatgtctg cacattggca ctttgacatg aattgatatc tcggcttgag actacctgng 300
tcaangctca catcagcccn gatgacgtgg tgtcccgaga canatgga 348

<210> 1610
 <211> 266
 <212> nucleic acid
 <213> Zea mays

 <400> 1610

 atcaagcccc agactgaccg ctgggtgttg atggtgatgc gcttgacgcc aggggnaggtc 60
 tcaaggccga gcatatcaat ttccattgtc aaagtggcca atgttgcaga caatggccag 120
 tgggtttcat ccttcctcat gtggtcaacc atgatgaata tccttgttgc cagtgggtgg 180
 tcacgaagat gtcagctttc cggaggnaaa cggctccann ctgtcgtga gaanncgtgt 240
 tgcnagnagg gtgcccgtgc cgcngc 266

<210> 1611
 <211> 143
 <212> nucleic acid
 <213> Zea mays

 <400> 1611

 gcgatacgg tgatgtcggc aagggttgtg ctgctgcact caaacaggct ggtgcccgtg 60
 tcattgtgac cgagatcnac cccatctgtg cctccaggc tctgatggag ggtcttcagg 120
 tcnttccctt ggagnacgtt ntt 143

<210> 1612
 <211> 118
 <212> nucleic acid
 <213> Zea mays

 <400> 1612

 aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca ctactccct 60
 gatgggtotga tgagggccac cgacgttatg atcgccggta aggttgccgt ggtctgcg 118

<210> 1613
 <211> 265
 <212> nucleic acid
 <213> Zea mays

 <400> 1613

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ctctgccgcc atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa 120
 ggattctcgc aggcggactt cggccgcctc gagattgagt ggccgaggtc gaaatgccgg 180
 ctcatggcgt gccgcggcga gttcggtccc tccaagcctt cggcggcgta agatctcggg 240
 gtctttcaca tgaacatcag accgc 265

<210> 1614
 <211> 111
 <212> nucleic acid
 <213> Zea mays
 <400> 1614

catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 60
 tgacaatgaa attgatatgc tcggccttga gacctaccct ggcgtcaagc g 111

<210> 1615
 <211> 154
 <212> nucleic acid
 <213> Zea mays
 <400> 1615

cnccagatc caattcgcga gtnncnccnc ctctgccgcc atggcgctct ctgtggagaa 60
 gacntcgtct ggacgggagt acaagggtcaa ggatctctcg caggcggacn tcggccgcct 120
 cgagattgag cnggccgagg tcgaaatncc cggc 154

<210> 1616
 <211> 226
 <212> nucleic acid
 <213> Zea mays
 <400> 1616

cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60
 cctcgtcctg gaacgggagn acaagggtcaa ggntntntng naggngnant tnggcngcnt 120
 cnanattnan ctggccnagg tcgaaatgcc cggcntnatg gcgtgcngcg ccganttcgg 180
 cccgtccaag ccttcgccgg cgcnaggntc tcgggggtctt cnacat 226

<210> 1617
 <211> 229
 <212> nucleic acid

<213> Zea mays

<400> 1617

ggatatcatc atggttgcca gtggtggtca cgaagatgtc agcttcagag acaacgtcct 60
ccaagggaag gacctgaaga cctccatca gagcctggag ggcacagatg gggtcganct 120
cggtcacaag gntatcatcn ngggtgacca catgaggag atgaagaaca atgccattgt 180
ctgcaacatt ggccactttg acaatgaaat tgatatgtc ggccttgag 229

<210> 1618

<211> 120

<212> nucleic acid

<213> Zea mays

<400> 1618

ctttgtontg tctgtctcat tcactaacca ggtcattgcc caacttgaac tgtggangga 60
gaagagctct ggcagatgtg anaagaaggt gtatgtgtc cccaagcacc ttgatgagaa 120

<210> 1619

<211> 109

<212> nucleic acid

<213> Zea mays

<400> 1619

gtgccctcca ggctctgatg gaggggtctc aggtccttcc cttggaggac gttgtctctg 60
aagctgacat cttcgtgacc accactggca acaaggatat catcatggt 109

<210> 1620

<211> 96

<212> nucleic acid

<213> Zea mays

<400> 1620

ggttgaccac atgaggaaga tgaagaacaa tgccattgtc tgcaacattg gccactttga 60
caatgaaatt gatatgtctg gccttgagac ctaccc 96

<210> 1621

<211> 118

<212> nucleic acid

<213> Zea mays

<400> 1621

cgctgagttc aagatcgtgc tcaccatcat ccgcgacggg ctcnaggctg accccaagaa 60
gtnccgcaag atgaaggaga ngcttgctcg cgtctctaag gagancancc angggtgt 118

<210> 1622

<211> 559

<212> nucleic acid

<213> Zea mays

<400> 1622

gnnnnnnngnn tnncnagann ntttcncttt ccaacnctc ttgaatttcc gggtcgaccc 60
acgcgtccgc aacctgtatg gntgccgnc a ctngatccct gatggtctga tgagggccac 120
tgaccgttat gntcgccgga aagggtgccc tggctcncgg atacngtgat ntcngcaagg 180
gttgtgctgc tgcaaatnaa gcanggctng tgcccgtnct attgtnaccc gagancnacn 240
natctgtncn nntacangct cttattngaa ggtctttang nccttcnctt ggaaganngt 300
ggctntgaag ctncatnttn ngaccaccac tgnaaacaag gatatnnnat ggttgaccac 360
atgangaana tgaagnacaa tgccattggc tgnaacattg ggccactttt gacaatgaan 420
ttgatatgct cnggccttga gacctaccct ggcgtcaaag cgcattatnc atcaaanccc 480
anactgaccg cttgggtgtt tcengagacc aaacactggc atcattggtc cttnctgaag 540
ggtcnnctgg ntnaacntt 559

<210> 1623

<211> 88

<212> nucleic acid

<213> Zea mays

<400> 1623

cgacaacgct gagttcaaga tcgtgctcac catcatccgc gacgggctca aggctgaccc 60
caagaagtac cgcaagatga aggagagg 88

<210> 1624

<211> 82

<212> nucleic acid

<213> Zea mays

<400> 1624

atctttctcca cgcaggacca cgcgcgcgcc gccatcgcg cgcactcggc cgcogtggtc 60
gcctggaagg gggagaccct cg 82

<210> 1625
<211> 139
<212> nucleic acid
<213> Zea mays

<400> 1625

cctcactccc gttccatttc cccatctccc agatccaatt cgcgagttct cctcctctg 60
ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120
tctcgcangc ggacttcgg 139

<210> 1626
<211> 255
<212> nucleic acid
<213> Zea mays

<400> 1626

agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc agcgtgccga 60
tcgagggtcc ctacaagcct gccactacc ggtactaggc acacggcttg cagctcactc 120
gggccgttgt gtgctatgaa gttcgctaca ctggcctgtc agttatcttt tgcattgcata 180
tgcattatca tatacgcagt cgcgtanagg ttttcttatg gttatcgctt gancngnngn 240
gggagggaag gagct 255

<210> 1627
<211> 224
<212> nucleic acid
<213> Zea mays

<400> 1627

acacngtccg ngacgctggg nactgtncng agaacgcgtn ggccggacgcg tgggtttccc 60
catatcccag atccanttcg cgagannctc cctcctangc ggccatgacg ctctctgtgg 120
agaagacctc gtctggacgg gagtacaagg tcaaggattc tcgcaggcgg acttcggccg 180
cctcgagatt gagtgccnag tgaatncccg gcnntgggg gngc 224

<210> 1628

<211> 113
 <212> nucleic acid
 <213> Zea mays

<400> 1628

cccgttccat ttcccatct cccagatcca attcgcgagt tctccctct ctgccgccat 60
 ggcgctctct gtggagaaga cctcgtctgg acgggagtag aaggtcaagg atc 113

<210> 1629
 <211> 182
 <212> nucleic acid
 <213> Zea mays

<400> 1629

tctcgttcc atttcccat gctccnagat ccaattccgc gagtncctcc ctccactgc 60
 ggccatggcg ctactctgtg gagaagacct cgtctggacc gggagtacca aggtcaagga 120
 tctctacgca ggcggaacttc ggccgntcga gattgagctg gccgaggtcg aaatgcccgg 180
 cc 182

<210> 1630
 <211> 107
 <212> nucleic acid
 <213> Zea mays

<400> 1630

ttccatttcc ccattctcca gatccaattc gcgagttctc cctcctctgc ngccatggcg 60
 ctctctgtgg agaagacctc gtctggacgg gagtacaaag ttaaagg 107

<210> 1631
 <211> 283
 <212> nucleic acid
 <213> Zea mays

<400> 1631

aaaatggctt ctacaggtccc tcacgttccg cctcgtgctg ctactgtcct ctcgcgaca 60
 gagtcatcg gaaatactcc tctcgttaga cttaacaaga tccccagtc gctgggcatc 120
 gaggcgatg tctacgtcaa gccagagctg ttcagcgctg gaggcagtgt taaggacaga 180
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cttatcgagc ctaccagtgg aaacactggg atcgggtcttg ctc 283

<210>	1632
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<400> 1632

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cqcgagggc gagagtggag tggatgatgcg ggtngccgtc gacgcgaanc gccagngtgg 120

gtgtagggcgc tcggatgctt ttctacccga cgctggtgta caacgtcgtg aggaatcggg 180

tcgagaagca cttccactgg tgggatacaga tcgatgagca tgtcctgctc ggtgctgttc 240

catccctagc gatgttctcc ggctaaaga 269

[illegible]

<210>	1633
<211>	125
<212>	nucleic acid
<213>	Zea mays

<400>	1633
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tinctggacng attgtcacat ggatnacnga tcnnatgcat ggaaacacca tcaaggcccc 60

ttgtggcctg aagacnccgt ccatttgacn ncatncnggc tgaagtnccg tncctncttc 120

gatgt 125

<210>	1634
<211>	123
<212>	nucleic acid
<213>	Zea mays

<400> 1634

gtgctgttcg caatgctgga ttaattgtca catggattac tgatcctatg catggaaaca 60

ccatcaaggc cccttgtggc ctgaagactc gtccattcga ctcaattctg gctgaagtgc 120

gcq 123

<210>	1635
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<212>	nucleic acid
<213>	Glycine max

<400> 1635

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gagaatggat accttctctgt tcaacttctga gtcggtgaac gagggtcac cgcacaagct 120
gtgtgatcag atatctgatg cagtgcctga tgcctgcctg gagcaagacc ctgaaagcaa 180
ggttgcctgt gagacatgca ccaagaccaa cttggatcatg gtatttggag agatcaccac 240
aaaagccaat atngactatg agaagattgt gcgcgatacg tgccgttcta taggatttgt 300
gtctggtgat gt 312

<210> 1636

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 1636

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ctgacaactg caaggctcctt gtaaaccattg agcagcagag ccctgatttg gccagggtgt 120
gcacggcaac ttaaccaaag acccgaggaa atcgggtgtg agaccagggt cactgtttgc 180
tatgcacgga cgagacccca gattgngcca tgagtcattg ncttncaata aactcgggtgt 240
cgtctaccga gntcgcaaga cggnaacttc cnggttgggc tatggaanac cna 293

<210> 1637

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 1637

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gtactacaat gacaagggtg ccatggttcc aatccgcgtc cactgtgtgc ncatctccac 120
acagcatgat gagnctgtc acaaatgatg ngattgcagc tganctnaaa gaacacgtga 180
ttaagcntgt gattccngng nagtaccttg nngagccgac cnntggccng tngaaccctc 240
tggcaggttt gncttgnagg ccgcgtgggg atgctggnt caccggcc 288

<210> 1638

<211> 292

<212> nucleic acid

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<213> Glycine max
 <400> 1638
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 ttanaatgac aatgggtgcc gggntcctat ncgngtacac acngtgctaa tntccaenca 180
 acacgangan nontgtccnc cantgacgan attgctgntg nctcaaaga gcatgtgntc 240
 aagnctgtgn ngcnagataa gtaccttggt gnggagncca tttncnttgn nc 292

<210> 1639
 <211> 285
 <212> nucleic acid
 <213> Glycine max
 <400> 1639
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 taaggccatt cattcctcac cgctgtgtgc tgggagtttt ttgagctttg ccottatcat 180
 atctataaatt tgtttcatatt attttactta attcgtgtgt gtttctcaact ttctctnoot 240
 cctctccatt ctattttggt tcttctatcc tcatatgtaa ttttt 285

<210> 1640
 <211> 275
 <212> nucleic acid
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 cgttctatag gatttgtgtc tggatgatgtt ggccttgatg ctgacaatgc aaggtattgg 120
 tgtaccttga gcagcaaagt cctgacatag cacaaggagt ccatggccac ctcaccaagc 180
 gaccagaaga cataggagct ggcgatcagg gtcacatgtt tggctacgcc acagatgaaa 240
 cacctgagct tatgcctctc agccatgtcc ttgca 275

<210> 1641
 <211> 317
 <212> nucleic acid

<213> Glycine max

<400> 1641

ttcannantn gcatgcacgc gtacgtaagc tngaattcn gctcganctc nagccgaatc 60
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ncttgnntgn gaagnccatt ttccagttca acccctctgg ccgttttgtc attngaggtc 180
ctcaaongtg atgctctctc caccagccga caagatccat tcnnccgnta cttanggagg 240
cnggggtgct catngtggtg gtgccttctc cgggaacgat cccaacaagg tttatangag 300
ttgtgttaca ntgtgag 317

<210> 1642

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 1642

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctctacaa ttttttttta 60
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gaagaaaacgc agaaactcat aaaccatcgc tgactccgca attttcgtcg gttcctcaca 180
cactctctga tttaancncg gtgccaagct caccgaggtt cggaagaacg ggacatgccc 240
ttggctgaga cctgatggca agaccaagt cactgttg 278

<210> 1643

<211> 259

<212> nucleic acid

<213> Glycine max

<400> 1643

gatctcaaga ggggtgggaa taacaggttc ttgaagactg ctgcatatgg acacttcggc 60
agagaggacc ctgacttcac atgggaagtg gtcaagcccc tcaagtggaa tgaatgggag 120
tttttttagcg ttgcccttat aatgtctntt atccataact ttccacgtcc cttgctctgt 180
gtttttctct cgtcgtctc ctcctatntt gtttctcng cctttcattt gtaatttttt 240
acatgatcaa ctaaaaaat 259

<210> 1644

<211> 191
 <212> nucleic acid
 <213> Glycine max

 <400> 1644

 gongcgtacg taagctcggn ntctggctcg aggagnattn tgaattcacg cctggaatga 60
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 cttatggaca ttttggaaga gatgatccgg atttcacatg ggagacagtg aagataactca 180
 agcctngtgc t 191

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 <400> 1647

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 tcccacaanc tgaactcggc catgaccaag gctcgcnagg attgtactct gccatggctg 180
 cgacccgaca ccaagactca ggtcactgtc gagtacgcc acgatggcgg tgccgtcgtc 240
 cctctccgtg ttgacaccgt cgtcgtctct gnccagcana ncgaggacat cactatcgag 300
 aagctccgcg aggagatcaa ggagaagatc at 332

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 gcagagccct gatattgcc aggggtgtgca cggccacett accaaaagac ccgaggaaat 180
 cgggtgctgga gaccagggtc acatgttttg ctatgccact gtcgagaccc cagaattgat 240
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 tggcctnaga aattncctat tcacnctnna tcagtgaacg ngnggcanc cgcacaggnn 180
 ctgtgaccag atctccgatg ctgtgntcng attcatgctt ggagcaggac cctgagagca 240

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<212> nucleic acid
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ctatncacat ctgantctgt anacgagggg caccocggac aagctgtgcy nccagatctc 180
tgatggcagt gctcgatgcy tgctgnaen ggacctgag cagcgagggt gcctgtnaga 240
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<400> 1651

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ccttcagtgg ntgtcttgct ggggtgtggat gaataatttg cgtgtttcat gactactact 180
actactactc cnttcnntgt ctaatgccat ctcatcnatn nctaaactgn tcgntttntt 240
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<212> nucleic acid
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 cttcggagaa atcacgacca aggccaatgt tgactacgag aagatagtgc gtgtacaccc 180
 tgagcaatat cagggctctg ctgctcaatg ttgacgagga ccttgagttt gtcggcatcc 240
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<210> 1653
 <211> 320
 <212> nucleic acid
 <213> Glycine max
 <400> 1653

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 ccttgaacag gaccctgaca gcaaggttgc ctgtgagaca tgcaccaaga ccaacatggt 180
 catgggtcttt ggagagatca caaccagggc aangtanann ntgagaagan tgtccnggan 240
 anangccgcg aaattggatt cnaccccgga nnanttggtt nnnnnnnna aaantnnngg 300
 ntnggncaan nnttggnnc 320

<210> 1654
 <211> 506
 <212> nucleic acid
 <213> Glycine max
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 cgtcagtacg gatgcgagaa gacgacagaa gggggcagcg cttgagacca agccccactc 120
 aaccaacaca acactctctc tgctctctct ctaactttca aagtttttta agtnnttaga 180
 tggcagagac attcctaatt acctcaaagt cncnaacga agganacctg acaagctctg 240
 cgaacaaatn tccgatgctg tncnnaanc ctgccttgaa caagacccaa acagcaaagt 300
 tncctgcaa acatgcccc agaccaactt tgcenangtc ctccgagaaa ttnccaccaa 360
 gggcaacntt nactacgana anatnntgcg ttacacctgc nggaacatcg ggttcntctc 420
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aaccctgaaa ntnnccaagg gggttg

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<210> 1655
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<212> nucleic acid
<213> Glycine max

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cttgagacca agccccactc aaccaccaca ccaactctctc tgctcntctt ctatctttca 180
acgttttttna agtattaaga tggcataaac attcctattt acttcatant cagtnaanga 240
gggacacnct gacgtinctc gcgannanct ctengattct gtcctcnacn cntgccttna 300
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cttcgggagag atnnccatna tggcnaacgt tgactancat gaagatcntg cntnacacct 420
gcaagaattc ggattcgtct caaaccatgt tggganntat gctgannntg naaggttcnt 480
tgnaaacggt tgnccntttc a 501

<210> 1656
<211> 533
<212> nucleic acid
<213> Glycine max

<400> 1656

angggtgata ttntgtgten aaaggatatag tangtgntag attntcttcg gtttcggtat 60
taccnacnng accacgcgt cccgcccacg cgtccggctg cgataagacg acagaanggg 120
atacctatgg tggctggggg gngggatggg ggaagtgcct tttgggggaa ngaccctacc 180
aaggttgaca gaagtgggtg ctatattgta aggcattgctg caaanagtgt ngtggcaaat 240
ggccttgcta naagggtgcat tgtgcaagtt tcctatncca ttggtgtccc tgaacccttg 300
tcanatgttt ntngacactt atngaactgn naanattcna nacaannagag attttinctat 360
anntatanga ataattttta nnttnananc tngnnnnngtn tnncataaan nttingtaant 420
aaataagnnn naantntttt gtnnatttaa naaattntan tnttnnttta natagnnaaa 480
taatttttna agtnnatttn nngtnnnntt nnaaattatg tannatatnt ctt 533

<400> 1657

[illegible]

<400>	1658
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ttcgggncga	cccacgcgnc	cggcnngccn	canngaacca	ccacacctct	tntcgttttt	120
gctacccent	tctgenctac	ggggaccggg	naagttttta	nngggtaang	atggcagaga	180
caatnnnatt	taccnnagag	tcggtgaacg	agggacaccc	tgacaagcnn	tgcgacaaaa	240
tncccnatgc	tgtcctcnac	gcttgcctng	agcaggaccc	anacagcaaa	gngtgnctgc	300
tgaaacatgc	acaaaaacca	actnggncat	gggcttgga	gaaatnacia	ccaaggncac	360
ggcngactac	gacaagatag	nncagagaca	cctgcangaa	cntnggacnt	cgnntnaaat	420
gaagtnggga	nttgatgcc	nnnaacttgc	caagntccnn	ngntaaccat	tgtacaatna	480
tgatccccgg	ngagttggtc	naggentaca	agggnccacn	ntntcnanaa	ancnggant	540
attngnnnn	tggtgnc					557

<210> 1659

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1659

cgtcgcangc acgcgtacgt aagctcggaa ttccggctcga gnnactttcc tcttcacctc 60
agaatctgta aacgaaggcc atcccgacaa gctgtgtgac caggtttcag atgccatcct 120
agatgcatgc ttggagcaag acccagaaaag caaggttgct tgcgagacct gtacaaaaac 180
taacatgggtt atgggtctttg gtgagattac aaccaaggcc agcgtgaact acgagaaaat 240
agttcgagac acttgcaaag gcattggggtt tgtgtcacca gatgt 285

<210> 1660

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 1660

cacgcgtacg taagctcgga attcggctcg agcttcctct tcgcacaaaag cagcaagcat 60
ccttgagatg gaaactttcc tcttcacctc agaatctgta aacgaaggcc atcccgacaa 120
gctgtgtgac caggtttcag atgccatcct agatgcatgc ttggagcaag acccagaaaag 180
caaggttgct tgcgagacct gtacaaaaac taacatgggtt atgggtctttg gtgagattac 240
aaccaaggcc agcgtgaact acgagaaaat agttcgagac acttgcaaag gcattggggtt 300
tgtg 304

<210> 1661

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 1661

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ntccatcttc ttottctctt 60
cctcttcgca caaagcagca agnatecttg agatggaaac ttctctnott cacctcagaa 120
tctgtaaacg aaggccatcc cgacaagctg tgtgaccagg tttcagatgc catcctagat 180
gcatgcttgg agcaagaccc agaaagcaag gttgcttgcg agacctgtac aaaaactaac 240
atggttatgg tctttgggtga gattacaacc aaggccagcg tga 283

<210> 1662

<211> 447

<212> nucleic acid

<213> Glycine max

<400> 1662

gagctggtga ccagggtcac atgtttgggt atgccaccga tgaaaccccc gagtacatgc 60
ccctcagcca tgtccttgca accaaacttg gtgctcgect cacagagggt aggaagaatg 120
gcacctgtgc ttggttgagg ccagatggta agacacaagt aaccgtcgag tactacaatg 180
acaatggtgc catggttcca gttcgtgtcc acactgtcct aatttccacc caacatgatg 240
agactgtgag caatgatcaa attgctgagg accttaaaga gcatgttatc aagcctgtca 300
ttcctgagaa gtaccttgat gagaagacca tcttccacct taacccttct ggccgttttg 360
tcattgggtg ccctcatggt gatgctggtc tcaactggaa gaaagatnat cattgatacc 420
taagggtggct ggggtgctca aggtgga 447

<210> 1663

<211> 475

<212> nucleic acid

<213> Glycine max

<400> 1663

ccacgcgtcc gcacaaagcg ggttactgtc tgttcaagct accatctctc tctctctttc 60
ttagngcctc cttgccagaa gttaaaatgg cccaagaaac tttcctattc acatctgaat 120
cagtgaacga ggggcacctt gacaagctct gtgaccagat ctccgatgct gtgctcgatg 180
catgcttgga gcaggacctt gacagcaagg ttgcctgtga aacctgcacc aagaccaaca 240
tggtgatggt tttcggagag atcacaacca aggccaacgt ggactatgag aagattgtgc 300
gtgacacatg caggaacatt ggttttgtct ctgatgatgt tggctttgat gctgacaact 360
gcaaggctct cgtcaacatt gagcaacaga gtccatgatat tgctcaaggt gtgcacggnc 420
acctnacaaa gaggcctgag gagattggtg ctggtgacca aggtcatatg ttccg 475

<210> 1664

<211> 520

<212> nucleic acid

<213> Glycine max

<400> 1664

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nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn gagaagacga cagaaggggt 120
ctcaaattgat gtgggactgg atgccgacaa ctgcaagggtc ctcgtcaaca ttgagcagca 180
gagccctgat attgctcagg gtgtacacgg ccaccttacc aaaaaacctg aagaaattgg 240
tgctggtgac cagggtcaca tgtttggtta tgccactgat gaaacctctg aattgatgcc 300
attgagccat gttcttgcaa caaaactcgg tgctcgtctc accgagggttc gcaagaacgg 360
tacctgccct tggctgaggg ctgatgggaa gacccaagtg accgttgagt attacaatga 420
caatggtgcc aggggttccta ttctgtgaca caccgtgcta atctccaccc aacacgacga 480
gactgtcacc aatgaccaa ttgntgntta acctnaaaaa 520

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<210>      1665
<211>      494
<212>      nucleic acid
<213>      Glycine max
<400>      1665

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ggngnnnnnga atgatctagg tnntctatgc cgagacatga cgnaagancc acngtacgt 60
aaactcgga ttcggtcga gaacagcaca aagcgggtta ctgtctgttc aagctaccat 120
ctctctctct ctttcttagt gctccttgc cagaagttaa aatggcccaa gaaactttcc 180
tattcacatc tgaatcagt aacgaggggc acctgacaa gctctgtgac cagatctccg 240
atgctgtgct cgatgcacgc ttggagcagg acctgacag caagggttgc tgtgaaacct 300
gcaccaagac caacatggtg atggttttcg gagagatnac aaccaaggcc aacgtggact 360
atgagaagat tgtgcgtgac acatgcagga acattggttt tgtctctgat gatgttggtc 420
ttgatgctga caactgcaag gtctcgtca acattgagca acagagtcct gatattgctc 480
aaggtgtgca cngg 494

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<210>      1666
<211>      502
<212>      nucleic acid
<213>      Glycine max
<400>      1666

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gnagtgtttg ntntgggggg ggggagnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnaaagcgg gttactgtct 120

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gttcaagcta ccattctctct ctctctttct tagtgcctcc ttgccagaag ttaaaatggc 180
ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccctg acaagctctg 240
tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg acagcaaggt 300
tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcggagaga tcacaaccaa 360
ggccaacgtg gactatgaga agattgtgcg tgacacatgc aggaacattg ggttttgtct 420
ctgatgatgt tgggtttgat gctgacaact gcaagggtccc tcgtcaacat tgagcaacag 480
agtcttgata ttgctcaagg tg 502

<210> 1667
<211> 372
<212> nucleic acid
<213> Glycine max
<400> 1667

1667
372
nucleic acid
Glycine max
1667

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acaagctctg tgaccagatn tccgatgctg tgctcgatgc atgcttggag caggacctga 120
cagcaagggtt gcctgtgaaa cctgnaccaa gaccaacatg gtgatgggtt tccgagagat 180
cacaaccaag gccaacgtgg actatgagaa gattgtgctg gacacatgca ggaacattgg 240
ttttgtctct gatgatgttg gtcttgatgc tgacaactgc aagtcctcgt caacattgag 300
caacagagtc ctgatattgc tcaagggtgtg cacggccact cacaaagagg cctgaggaga 360
ttggtgtggt na 372

<210> 1668
<211> 487
<212> nucleic acid
<213> Glycine max
<400> 1668

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ttcggctcga ggtcgggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat 120
gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 180
acaaaaacca acttgggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 240
gagaagatag tgcgtgacac ctgcaggaac atcggttcg tctcaaatga tgtgggactg 300

gatgccgaca actgcaaggt cctccgtcaa cattgagcag cagagccctg atattgctca 360
 aggtgtacac gggcaactta ccaaaaaaacc tgaagaaatt ngtgctggtg accaaggtca 420
 cattttgggt aatnccactg gntgnaaacc ccngnattta tgncccattg accnagttct 480
 tnncaaa 487

<210> 1669
 <211> 419
 <212> nucleic acid
 <213> Glycine max
 <400> 1669

Sequence 1669-419

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 tctctctctc tttcttagtg cctccttgcc agaagttaaa atggccaag aaactttcct 120
 attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc agatctccga 180
 tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggttgct gtgaaacctg 240
 caccaagacc aacatggtga tggttttcgg agagatcaca accaaggcca acgtggacta 300
 tgagaagatt gtgcgtgaca catgcaggaa cattggtttt gtctctgatg atgttgggtc 360
 ttgatgctga caactgcaag gtctctngtca acattgagca acagaatcct gatattgct 419

<210> 1670
 <211> 447
 <212> nucleic acid
 <213> Glycine max
 <400> 1670

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 tcttcagcng gngaaatggc acangaaacc tttctattca catctgaatc tgnaaacgag 120
 ggtcaccocg acaagctgtg cgaccagatc tctgatgcag ngctcgatgc gtgccttgaa 180
 caggaccctg acagcaaggt tgctgtgag acatgcacca agaccaacat ggtcatggtc 240
 tttggagaga tcacaaccaa ggccaacgta gactatgaga agattgtccg tgacacatgc 300
 cgngaaattg gattcatctc tgatgatgtt ggtcttgatg ctgacaaatg caaggngttg 360
 gtcaacattg aacancaaan ccctgatatc nccaggngn gcacggnac ttcaccaacc 420
 cccaaaagaa ggttnggctn ggncca 447

<400> 1671

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1970	1971	1972	1973	1974	1975	1976	1977	1978	1979	1980	1981	1982	1983	1984	1985	1986	1987	1988	1989	1990	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099	2100	

<400>	1672
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tgcaagaag	acgacagaag	ggggcacccgc	ttgagcagac	ttaacaacag	cacaaagcgg	120
gttactgtct	gttcaagcta	ccatctctct	ctctctttct	tagtgccctcc	ttgccagaag	180
ttaaaatggc	ccaagaaact	ttcctattca	catctgaatc	agtgaacgag	gggcaccctg	240
acaagctctg	tgaccagatc	tccgatgctg	tgctcgggtgc	atgcttgag	caggaccctg	300
acagcaaggt	tgccctgtgaa	acctgcacca	agaccaacat	ggtgatgggt	ttcggagaga	360
tcacaaccaa	ggccaacgtg	gactatgaga	agattgtgcg	tgacacatgc	aggaacattg	420
gttttgtctc	tgatgatggt	ggtcttgatg	ctgacaactg	caaaggctct	cgtcaacatt	480
gagcaacaaa	at					492

<210>	1673
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<211> 503
 <212> nucleic acid
 <213> Glycine max

<400> 1673

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 cnaagaccaa agccccactc aaacaacaca ccaatctctc tgctcctcct cnaactttca 120
 agttttttaa gtnttaaaga tggcagagac attcctaatt nacctcagag tcagtgaacg 180
 agggacaccc tgacaagctc tgcgaccaaa tctccgatgc tgtcctcnac gcttgccctg 240
 aacaggaccc agacagcaan gttgcctgcg aaacatgcac caagaccaac ttggatcatg 300
 tcttcggaga gatnaccanc aaggccaacg ttgactacga gaagatngtg cgtgacacct 360
 gcangaacat cggttcctgc tcaaacgatg tgggacttga tgctgacaac tgcaaggctc 420
 ttgtaaacat tgagcaacaa aaccctgata ttgcccaagg tgtncacggc caacttacca 480
 aaaganccga aggaaatcng tgc 503

<210> 1674
 <211> 508
 <212> nucleic acid
 <213> Glycine max

<400> 1674

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 ntacgtaagc tcggaattcg gctcgagggtg gactatgaga agattgtgcg tgacacatgc 120
 aggaacattg gttttgtctc tgatgatgtt ggtcttgatg ctgacaactg caaggctcctc 180
 gtcaacattg agcaacagag tcttgatatt gctcaagggtg tgcacggcca cctcaciaag 240
 aggcctgagg agattggtgc tggtgaccaa ggtcatatgt tcggctatgc cactgacgag 300
 actcccgagc tcatgccctt gagccatgtc cttgccacga agctcgggtc caagctcanc 360
 gacgggtccg aaaaacngga aatgcccttg ggctgaaaac ctgatggcaa nnaccaagtc 420
 actgttggnn tactacaatt gacaaggggt ccatgggtcc aatccgcgtc aaaactgttg 480
 ctcatntcca anacagcaat gatngaga 508

<210> 1675
 <211> 334
 <212> nucleic acid

<213> Glycine max

<400> 1675

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gctgcaaaga gtgtcgtggc aaatggcctt gctagaagggt gcattgtgca agtttcctat 120
gccattgggtg tccctgagcc cttgtcagtgtttgtggaca cttatggaac tgggaagatt 180
cctgacaagg agattctgca aattgtgaag gagaatttcg acttcagacc tggaatgatc 240
accattaact tggaccttaa gaggggtggt catagggtcc tcaagacagc tgcttatgga 300
cactttggaa gggatgatgc agacttcacc tggg 334

<210> 1676

<211> 335

<212> nucleic acid

<213> Glycine max

<400> 1676

agtcgcatgc acgcgtacgt aagctcgga ttcggctcga ggcaagggtt cctgtgaaac 60
ctgcaccaag accaacaatgg tgatggtttt cggagagatc acaaccaagg ccaacgtgga 120
ctatganaag attgtgcgtg acacatgcag gaacattggt nttgtctctg atgatgttgg 180
tcttgatgct gacaactgca aggtcctcgt caacattgag caacagagtc ctgatattgc 240
tcaagggtgtg caaggccacc tcacaaagag gcctgnggag attggtgctg gtgaccaagg 300
tcatatgttc ggotatgcca ctgacgagac tcccg 335

<210> 1677

<211> 337

<212> nucleic acid

<213> Glycine max

<400> 1677

aaagatgcct gaggagattg gtgctggtga ccaagggtcat atgttcgggt atgccactga 60
cgagactccc gatctcatnn cttgagccat gtcccttgcca cgaagctcgg ngccaagctc 120
accgaggttc gnaagaacgg gacatgcctt tggctgagac ctgatggcaa gaccaagtc 180
actgttgagt actacaatga caagggtgcc atggttccaa tccgcgtcca cactgtgctc 240
atctccacac agcatgatga gccctgtcac aaatgatgag attgcagctg atcttaaaga 300

acacgtgant aagcctgtga ttccctgagaa gtacctt

337

<210> 1678
<211> 448
<212> nucleic acid
<213> Glycine max

<400> 1678

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gcaagtgtct tatgccattg gtgtgcctga gcctttgtct gtgtttgttg acacctatgg 120
cactgggaag atccatgata aggagattct caacattgtg aaggaaaact ttgatttcag 180
gcctggtatg atctccatca accttgatct caagaggggt ggaaataaca ggtttttgaa 240
gactgctgcc tatggacact ttggaagaga agaccctgac ttcacatggg aagtggtaa 300
accctcaag tgggagaagg cctaagtaat tcattccact gctctatgct ggaagttttt 360
tgagcgttgc cttataata tgtctaata ccataacttt ccacgtctct tactctgtgt 420
gtttctctcc tnttctctta ttttgga 448

CGCGTCCGCCACGCCTCGCGCCACGC

<210> 1679
<211> 336
<212> nucleic acid
<213> Glycine max

<400> 1679

tgcangcgta cgtaagctcg gaattcggct cgagncgtct aagcagcatt gtggcaagtg 60
gacttgccag aaggtgcatt gtgcaagtgt cttatgcat tgggtgcct gagcctttgt 120
ctgtgtttgt tgacacctat ggcactggga agatccatga taaggagatt ctcaacattg 180
tgaaggaaaa ctttgatttc aggcctggta tgatctccat caaccttgat ctcaagaggg 240
gtggaaataa cagggtttttg aagactgctg cctatggaca ctttggaaga gaagacctg 300
acttcacatg ggaagtggtc aaacctca antggg 336

<210> 1680
<211> 493
<212> nucleic acid
<213> Glycine max

<400> 1680

aggtcctcgt caacattgag caacagagtc ctgatattgc tcaaggtgtg cacggccacc 300
tcacaaagag gcctgag 317

<210> 1683
<211> 406
<212> nucleic acid
<213> Glycine max
<400> 1683

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atttacctca gagtcagtga acgagggaca ccctgacaag ctctgcgacc aaatctccga 180
tgctgtcctc gacgcttgcc ttgaacagga cccagacagc aaggttgcct gcgaaacatg 240
caccaagacc aacttggtca tgggtcttcgg agagatcacc accaaggcca acgttgacta 300
cgagaagatc gtgctgacga cctgcaggaa catcggttc gtctcaaacg atgtgggact 360
tgatgctgac aactgcaagg tccttggtta cattgagcaa caaaag 406

<210> 1684
<211> 489
<212> nucleic acid
<213> Glycine max
<400> 1684

actccaccgc gncggtaccg ttntaagncc ccgggccgac aaacgcgtca gtccggctgc 60
gagaagacga cagaaggggc accgcttgag cagacttaac aacagcaca agcgggttac 120
tgtctgttca agctaccatc tctctctctc tttcttagtg ccttcttgcc agaagttaaa 180
atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 240
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aaggttgcct gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagatnaca 360
accaagggca acgtggacta tgagaagatt gtgctgacga catgcaagaa cattgggttt 420
gtctctgatg aagttgggtc tgatgctgac aactgcaang gtctcgttca acattgcagc 480
cancaagat 489

<210> 1685

<211> 506
 <212> nucleic acid
 <213> Glycine max

<400> 1685

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 gcgtcaagta cggtgcgcnag aagacgacag aaggggatga aaccctgag tacatgcccc 120
 tcagccatgt ccttgcaacc aaactcgggtg ctgcctcac cgaggtagg aaaaatggta 180
 cctgtgcttg gctgaggcca gatggcaaga cacaagtaac tggtgagtac tacaatgaca 240
 atggtgccat ggttccagtt cgtgtccaca ctgtcctaata ttccacccaa catgttgaga 300
 ctgtgagcaa tgaccaaatt gctgctgacc ttaaagaaca tggtatcaag cctgtcattc 360
 ctgagaagtn cctggatgag aagaccatct tccaacctta aaccttctgg gcgtttttgn 420
 cnnttggtgg gcccnnangg tganncccgg gcccanatgg gaaannaaag atttccccnt 480
 ggaaaccocan aggttggnngn gggntc 506

<210> 1686
 <211> 427
 <212> nucleic acid
 <213> Glycine max

<400> 1686

gaggccagge aagccccact caaccaccac acctctctc gggtcacgcc taccctttct 60
 gctctttcttc nacctttcaa gttttaaaag tataaagatg gcanagacat tcttatttac 120
 ctgagagtgc gtgaacgagg gacacctga caagctctgc gaacaaatct ccgatgctgt 180
 cctcgacgct tgctcgagc aggaccana cagcaaagtt gcctgcgaaa catgcaccaa 240
 aaccaacttg gtcatggtct tcggagaaat cagaccaag gccaacgttg actacgagaa 300
 gatagtgcgt gacacctgca ggaacatcgg cttcgtctca aatgatgtgg gactgggatg 360
 ccgacaactg caaggtctc gtcaacattg agcancagan ccctgatatt gccanggtg 420
 tacacgg 427

<210> 1687
 <211> 504
 <212> nucleic acid
 <213> Glycine max

<400> 1687

ggnaactctt cgcggccaaa ctcttacann nccaggtagn gntanangaa ttcccggctc 60
gacccacgcg tnacgtacgg ctgcgagaag acgacagaag ggggcagcgc ttgagaccaa 120
gccccactca accaccacac cactctctct gctcttcttc tacctttcaa gtttttaaag 180
tattaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 240
caagctctgc gaccaaattc cccatgctgt cctcgacgct tgccttgaac aggaccaga 300
cagcaagggt gcctgcgaaa catgcaccaa gaccaacttg gtcattgtct tcggagagat 360
caccaccaag gccaacgttg actacganga gatcgtgcgt gacacctgca ggaacatcgg 420
cttcgtctca aacgatgtgg gacttgatgc tgacaactgc aaggctcctg taaacattga 480
agcagcagag ccctgatatt gcc 504

<210> 1688

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 1688

ncgnangcac gcgtacgtna gctcggaatt cggctcgagn ctgaatcngt gaacgagggg 60
caccctganc aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagca 120
ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaactatg tgatgggttt 180
cggagagatc acaaccaagg ccaacgtgga ctatgagaag attgtgcgtg acacatgcag 240
gaacattggg tttgtctctg atgatgttg tcttgatgct gacaactgca aggtcctcnt 300
caacattgag caacagagtc ctg 323

<210> 1689

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 1689

tcggctcgag ngaccatctt ccaccttaac ccttctggcc gttttgtcat tgggtggcct 60
catgggtgatg ctgggtctcac tggaagaaa atcatcattg atacctatgg tggctggggg 120
gctcatgggt gaggtgcctt ttcagggaa gaccctacca aggttgacag aagtgggtgcc 180

tatattgtaa ggcaggctgc aaagagtgtc gtggcaaattg gccttgctag aaggtgcatt 240
 gtgcaagttt cctatgccat tgggtgtccct gagcccttgt cagtgtttgt ggacac 296

<210> 1690
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 1690

gcacgcgtac gtaagctcgg aattcggctc gagtatgaga agattgtccg tgacacatgc 60
 cgcgaaattg gattcatctc tgatgatgtt ggtcttgatg ctgacaaatg caaggtgttg 120
 gtcaacattg agcagcagag ccttgatata gcccaggggtg tgcacgggtca cttaccaag 180
 cgcccagagg aggttggtgc tgggtgaccag ggtcacatgt ttggctatgc cactgatgaa 240
 acccctgagt acatgcccct cagccatgtc cttgcaacca aactcgggtgc tcgcctcacc 300
 gag 303

<210> 1691
 <211> 336
 <212> nucleic acid
 <213> Glycine max
 <400> 1691

gncnatccaa agcgtacgta agctcggaat tcggctcgag ggaacatcgg ctctgtctca 60
 aatgatgtgg gactggatgc cgacaactgc aaggctcctcg tcaacattga gcagcagagc 120
 cctgncattg ctacgggtgt acacggccac cttacaaaaa aacctgaaga aattgggtgct 180
 ggtgaccagg gtcacatgtt tggctatgcc actgatgaaa cccctgaatt gatgccattg 240
 agccatgttc ntgcaacaaa actcgggtgct cgtctcaccg aggttcgcaa gaacggtacc 300
 tgcccttggc tgaggcctga tgggaagacc caagtg 336

<210> 1692
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 1692

tcgcatgcnc gcgtacgtna gctcgggaatt cggtcggagg ttaaaatggc ccaaganact 60

ttcctattca catctgaatc agtgaacgag gggcacctcg acaagctctg tgaccagatc 120
 tccgatgctg tgctcgatgc atgcttggag caggaccctg acagcaagggt tgctgtgaa 180
 acctgcacca agaccaacat ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg 240
 gactatgaga agattgtgcg tgacacatgc aggaacattg gttttgtctc tgatgatgtt 300
 ggtcttgatg ctga 314

<210> 1693
 <211> 321
 <212> nucleic acid
 <213> Glycine max
 <400> 1693

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tcattatcga tacttatgga 60
 ggatgggggtg ctcatgggtg tgggtgctttc tccgggaagg accctaccaa ggttgatagg 120
 agtgggtgctt acattgtgag acaggctgct aagagcattg tggcaagtgg acttgccaga 180
 aggtgcattg tgcaagtgtc ttatgccatt ggtgtgcctg agcctttgtc tgtgtttgtt 240
 gacacctatg gcaactgggaa gatccatgat aaggagattc tcaacattgt gaaggaaaac 300
 tttgatttca ggcttggtat a 321

<210> 1694
 <211> 514
 <212> nucleic acid
 <213> Glycine max
 <400> 1694

gngnnnnagt ggntngtgna gtnnactnag nnaaatTTTtg naccggtcog gaattccogg 60
 gtcgaccac gcgtccgtac gagaagacga cagaaggggg cagcgcttga tttgaggcca 120
 ggcaagcccc actcaaccac cacacctctc ctcgttcagc ctaccccttt ctgctcttct 180
 tctacctttc aagtttttaa agtataaaga tggcagagac attcctatit acctcagagt 240
 cgggtgaacga gggacaccct gacaagctct gcgaccaaT ctccgatgct gtccctcgacg 300
 cttgcctcga gcaggaccca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 360
 tgggtcatggt cttcggagaa atcacgacca aggccaacgt tgactacgag aagatagtgc 420
 gtgacacctg caggaacatc ggcttcgtct caaatgatgt gggactggat gccgacaact 480

aactcttacg tngccaggnn ccggtanaga attaccgggg ncgacccacg cgtcngccca 60
cncgtccgcc cagcggtccg acgggtgcga gaagacgaca gaaggggggc agcgcttgag 120
accaagcccc actcaaccac cacaccactc tctctgctct tcttctacct ttcaagtttt 180
taaagtatta agatggcaga gacattccta ttacctcag agtcagtga cgagggacac 240
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgcc tgaacaggac 300
ccagacagca aggttgccctg cgaaacatgc accaagacca acttggtcat ggtcttcgga 360
gagatnacca ccaaggccaa cgttgactac gagaagatcg tgcgtgacac ctgcaggaac 420
atcggttcg tctcaaacga tgtgggactg atgctgacaa ctgcaaagtc cttgttaaca 480
atgaacanca aancoc 496

1698
300
nucleic acid
Glycine max
1698
60
120
180
240
300

<210> 1698
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 1698
cgcatgcacg cgtacgttag ctcggaattc ggctcgagg agtgggtgcct acattgtgag 60
gcaagctgca aagagcattg ttgcaaattg acttgctagg agggcaattg tgcaagtttc 120
ctatgccatt ggtgtgcctg agcccttgtc tgtgtttgtt gacacttatg gcactgggaa 180
gatccctgac aaggaaatcc tcagcattgt gaaggagagt tttgaacttca ggccctggcat 240
gatctccatc aaccttgatc tcaagagggg tggaaatggc aggttccttga agactgctgc 300

<210> 1699
<211> 303
<212> nucleic acid
<213> Glycine max
<400> 1699

acntnctgca cgcgtacgta agctcggaat tcggctcgag gtcatggtct ttggagagat 60
cacaaccaag gccaacgtag actatgagaa gattgtccgt gacacatgcc gcgaaattgg 120
attcatctct gatgatgttg gtcttgatgc tgacaaatgc aaggtgttgg toaacattga 180
gcagcagagc cctgatatcg ccagggtgt gcacggtcac ttcaccaage gccagagga 240
ggttggtgct ggtgaccagg gtcacatggt tggctatgcc actgatgaaa cccctgagta 300

cat

303

<210> 1700
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 1700

ncgcnnngcac gcntacgtna gctcggaatt cggctcgacg caccaagacc aacatggtga 60
tggttttcgg agagatcaca accaaggcca acgtggacta tgagaagatt gtgcgtgaca 120
catgcaggaa catttggtntt gtctctgatg atgttggtct tgatgctgac aactgcaagg 180
tcctcgtaa cattgagcna cagagtcctg atattgctca aggtgtgcac gnccacctca 240
caaagaggcc tgaggagatt ggtgctggtg accaaggta tatgttcggc tatgccactg 300
acgagactcc c 311

<210> 1701
<211> 425
<212> nucleic acid
<213> Glycine max

<400> 1701

gagtaacaac agcacaaagc gggttactgt ctgttcaagc taccatctct ctctctcttt 60
cttagtgccct ccttgccaga agttaaaatg gcccaagaaa ctttcctatt cacatctgaa 120
tcagtgaacg agggggcacc tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat 180
gcatgcttgg agcaggaccc tgacagcaag gttgcctgtg aaacctgcac caagaccaac 240
atggtgatgg ttttcggaga gatcacaacc aaggccaacg tggactatga gaagattgtg 300
cgtgacacat gcaggaacat tggttttgtc ccgatgatgt ttggtcctga tgctgacaac 360
tgcaangtcc cccgtcaaca atgagcaaca nagtcctga aaattgcna angngttgna 420
cgggc 425

<210> 1702
<211> 321
<212> nucleic acid
<213> Glycine max

<400> 1702

tgcnccgcta cgtaagctcg gnatttnggc tcgaagcatt gtggcaagtg gactagccag 60
aaggtgcatt gtgcaagtnt cttatgccat tgggtgtgcc gagcctttgt ctgtctttnt 120
tgacacctat ggcaccgggn agatccatga taaggagatt ctnacattg tgaaggagaa 180
ctttgatttc aggcccggta tgatctccat caaccttgat ctcaagaggg gtgggnataa 240
caggttcttg aagactgctg catatggnga cttcggcaga gaggaccctg acttcacatg 300
ggaagtggtc nagccctca a 321

<210> 1703
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 1703
tcgcangcac gcntacgtaa gctcggatt cggctcgagc ggctcgaggt agactatgag 60
aagattgtcc gtgacacatg ccgcgaaatt ggattcatct ctgatgatgt tggctcttgat 120
gctgacaaat gcaaggtggt ggtcaacatt gagcagcaga gccctgatat cggccagggt 180
gtgcacggtc acttcaccaa gcgcccagag gaggttgggt ctggtgacca gggtcacatg 240
tttggtatg ccaactgatga aaccctgag tacatgcccc tcagccatgt ccttgcgccc 300
aaactcgggtg n 311

<210> 1704
<211> 473
<212> nucleic acid
<213> Glycine max

<400> 1704
ttaacttgen cgcgccaggt ancggtcaag gaattcccg gtcgaccac ggcgcgagc 60
gctgcgagaa gacgacagaa gggggcagcg cttganacca agccccactc aaccaccaca 120
ccactctctc tgctcttctt ctaccttca agtttttaaa gtantaagat ggcagagaca 180
ttctatttta cctcanagtc agtgaacgag ggacaccctg acaagctctg cgaccaaata 240
tcogatgctg tctcgcagc ttgccttgaa caggaccag acagcaaggt tgctgcgaa 300
acatgcacca ngaccanctt ggtcatgggt cttcggagag atcaccacca aggccaacgt 360
tgactacgag aagatcgtgc gtgacacctg caggaacatc ggcttcgtct caaacgatgt 420

gggacttgat gctgacaact gcaaggtcct tgtaaacatt gagcagcaga gcc 473

<210> 1705
 <211> 319
 <212> nucleic acid
 <213> Glycine max
 <400> 1705

gcacgcgtac gtaagctcgg aattcggctc gagcaagtaa ctgttgagta ctacaatgac 60
 aatgggtgcc aatgttccagt tcgtgtccac actgtcctaa tttccacca acatgatgat 120
 tctgtgagca atgaccaa atgctgtgac cttaaagagc atgttatcaa gcctgtcatt 180
 cctgagaagt acctggatga gaagaccatc ttccaacctt aacctttctg gccgttttgt 240
 cattgggtggc cctcatgggtg atgctgggtct cactggaaga aagatcatca ttgataccta 300
 tgggtgggtgg ggtgctcat 319

<210> 1706
 <211> 507
 <212> nucleic acid
 <213> Glycine max
 <400> 1706

gnnnnnnaan tctacgccgc ccctctaacg ngcacaanat tcncgggaac gacccaacgcg 60
 nccgtacggc tgccaagaag acgacagaag ggggcagcgc ttgagancaa nccccactca 120
 accaccacac cactctctct gctcttcttc nanttttcaa gtttttaaag tattaagatg 180
 gcagagacat tctatattac ctgagagtca gtgaacgagg gacaccctga caagctctgc 240
 gaccaaattc ccgatgctgt cctcgacgct tgcctttaaa cangacccaa gacagcaaag 300
 ttgcctgcca aacatgcacc aagaccaact tggatcatggc ctccggagag atnaccacca 360
 agggcaacgt tgactacgag aagatcgtgc gtgacacctg caggaacatc ggcttcgtct 420
 caaacgatgt gggacttgat gctgacaact gcaangtcct tgtaaacatt gagcaacaaa 480
 acctganaa tncccaaggt ttcaccg 507

<210> 1707
 <211> 351
 <212> nucleic acid
 <213> Glycine max

<400> 1707

ccttttcagg gaaggaccct accaagggtg acagaagtgg tgcctatatt gtaaggcagg 60
ctgcaaagag tgtcgtggca aatggccttg ctagaagggtg cattgtgcaa gtttcctatg 120
ccattgggtgt ccctgagccc ttgtcagtgt ttgtggacac ttatggaact gggaagattc 180
ctgacaagga gattctgcaa attgtgaagg agaatttcga cttcagacct ggaatgatca 240
ccattaactt ggaccttaag aggggtgggc atagggtcct caagacagtg cttatggaca 300
ctttggaagg gatgatgcag cttcactggg aagtgtgaac cactcaagtc a 351

<210> 1708

<211> 509

<212> nucleic acid

<213> Glycine max

<400> 1708

ggngnnnnng gnnnnnnngn naacttttac gcngtccgt gccggtcana gaattcacgg 60
gnogacncac gcgtccnccc acgcgtccgc ccacgcgtcc gccacgcgt ccgctgcgag 120
aagacgacag aagggggcag cgcttgagac caagccccac tcaaccacca caccactctc 180
tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga cattcctatt 240
tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgacaaaa tctccgatgc 300
tgtcctcgac gcttgccctg aacaggaccc agacagcaag gttgcctgcg aaacatgcac 360
caagaccaac ttggatcatg tcttcggaga gatcaccacc aaggccaacg ttgactacga 420
gaagatcgtg cgtgacacct gcaggaacat cggcttcgtc tcaaacgatg tgggacttga 480
tgttgacaac tgcaaggctc ttgtnaaca 509

<210> 1709

<211> 267

<212> nucleic acid

<213> Glycine max

<400> 1709

gagacaggct gctaagagca ttgtggcaag tggacttgcc agaagggtgca ttgtgcaagt 60
gtcttatgcc attggtgtgc ctgagccttt gtctgtgttt gttgacacct atggcactgg 120
gaagatccat gataaggaga ttctcaacat tgtgaaggaa aactttgatt tcaggcctgg 180

tatgatctcc atcaaccttg atctcaagag ggggtggaaat aacaggtttt tgaagactgc 240
 tgcctatgga cactttggaa gagaaga 267

<210> 1710
 <211> 320
 <212> nucleic acid
 <213> Glycine max
 <400> 1710

acgtgcgacg cagcgcgtacg taagctcgga attcggctcg aggcagcaga gccctgatat 60
 cgcccagggt gtgcacggtc acttcaccaa gcgccagag gaggttggtg ctggtgaccn 120
 cggtcacatg tttggctatg ccactgatga aaccctgag tacatgcccc tcagccatgt 180
 ccttgcaacc aaactcgggtg ctgcctcac cgaggtagg aaaaatggta cctgtgcttg 240
 gctgaggcca gatggcaaga cacaagtaac tgttgagtac tacaatgaca atggtgccat 300
 ggttccagtt cgtgtccaca 320

<210> 1711
 <211> 330
 <212> nucleic acid
 <213> Glycine max
 <400> 1711

nnnaaaannt gacntcgcan gcacgcgtac gtaagctcgg aattcggctc gagggactgg 60
 atgccgacaa ctgcaaggtc ctgctcaaca ttgagcagca gagccctgat attgctcagg 120
 gtgtacacgg ccaccttacc aaaaaacctg aagaaattgg tgctgggtgac cagggtcaca 180
 tgtttggcta tgccactgat gaaaccctg aattgatgcc attgagccat gttcttgcaa 240
 caaaactcgg tgctcgtctc accgaggttc gcaagaacgg tacctgccct tggtgagge 300
 ctgatgggaa gaccaagtg accgttgagt 330

<210> 1712
 <211> 313
 <212> nucleic acid
 <213> Glycine max
 <400> 1712

agtngcangc acgcgtacgt aagctcggaa ttcggctcga gtgcaagttt octatgccat 60

tgggtgtgcct gagcccttgt ctgtgtttgt tgacacttat ggcactggga agatccctga 120
 caaggaaatc ctcagcattg tgaaggagag ttttgacttc aggcctggca tgatctccat 180
 caaccttgat ctcaagaggg gtggaaatgg caggttcttg aagactgctg catatggaca 240
 ctttggcaga gatgaccctg acttcacatg ggaagtgggtg aagccactca agggggagaa 300
 ggtacctgct tac 313

<210> 1713
 <211> 486
 <212> nucleic acid
 <213> Glycine max
 <400> 1713

gtttaactttc ccgcgcncgg tcangaataa acgggtcgan ccacgcgtcc gacggatgcg 60
 agaagacgac agaagggggc agcgcttgat ttgaggccag gcaagcccca ctcaaccacc 120
 acacctctcc tcgttcaogc taccoccttc tgctcttctt ctacctttca agtttttaaaa 180
 gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 240
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 tcacgaccaa ggccaacgtt gactacgaga agatagtgcg tgacacctgc angaacatcg 420
 gcttcgtctc aaatgatgtg ggactggatg ccgacaactg caaggtcctc gtcaacattg 480
 agcaac 486

<210> 1714
 <211> 474
 <212> nucleic acid
 <213> Glycine max
 <400> 1714

aactttacgc tgccangtnc cggtcnaga attnacgggg ccgaccacg cgtccntacg 60
 gctgcgagaa gacgacagaa gggggcagcg cttgagacca agcccaactc aaccaccaca 120
 ccactctctc tgctcttctt ctacctttca agtttttaaa gtattaagat ggcagagaca 180
 ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg cgaccaaatc 240
 tccgatgctg tctcgcagc ttgccttgaa caggaccag acagcaaggt tgcttgcgaa 300

acatgcacca agaccaactt ggtcatggtc ttcggagaga tcaccaccaa ggccaacggt 360
gactacgana agatcgtgcg tgacacctgc aggaacatcg gcttcgtctc aaacgatgtg 420
ggacttgatg ctgacaactg caaaggctct ttgtaaacad tgagcaacan agnc 474

<210> 1715
<211> 382
<212> nucleic acid
<213> Glycine max
<400> 1715

gtcgcangca cgcgtacgta agctcgggaa ttcgggctcg agcaacagca caaagcgggt 60
tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180
aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac 240
agcaagggtg cctgtgaaac ctgcaccaag accaacadtg tgatgggtttt cggagagatc 300
acaaccaagg ccaacgtgga ctatgagaag attgtgctgt acacatgcag gaacatttgt 360
tttgtctctg atgatgttgg tc 382

<210> 1716
<211> 308
<212> nucleic acid
<213> Glycine max
<400> 1716

nntcgcangc acgcgtacgt aagctcggaa ttcggctcga gggcaccctg nacaagctct 60
gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 120
ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca 180
aggccaacgt ggactatgag aagattgtgc gtgacacatg caggaacatt ggttttgtct 240
ctgatgatgt tggctcttgat gctgacaact gcaaggctct cgtcaacatt gagcaacaga 300
gtcctgat 308

<210> 1717
<211> 312
<212> nucleic acid
<213> Glycine max

<400> 1717

gtcnnangca cgcgtacgta agctcggaat tcggctcgag ccagatctcc gatgctgtgc 60
tcgatgcatg cttggagcag gaccctgaca gcaagggtgc ctgtgaaacc tgcaccaaga 120
ccaacatggt gatggttttc ggagagnnca caaccaaggc caacgtggac tatgagaaga 180
ttgtgctgta cacatgcagg aacattggtt ttgtcncatga tgatgttggt cttgatgctg 240
acaactgcaa ggtcctcgtc aacattgagc aacagagtcc tgatattgct caagggtgtgc 300
cgccacctc ac 312

<210> 1718

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 1718

gtcgcangca cgcgtacgta agctcgggaa ttccggctcga ggcgctggtg accaggggtca 60
catgtttggc tatgccactg atgaaacccc agaattcatg ccattgagtc atgttcttgc 120
aaccaagctc ggtgctcgtc tcaccgaggt tcgcaagaac ggaacctgcc catggctgag 180
gcctgatggg aagacccaag tgactgtgga gtattacaat gataatggtg ccaggggtcc 240
agttcgtggt cacaccgtgc taatctccac ccagcatgat gagactgtca ccaacgacga 300
aattgoggct gacct 315

<210> 1719

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 1719

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<213> Glycine max

<400> 1733

ngtcgcangc acgcgtacgt aagctcggaa ttcggctcga gattgatacc tatgggtgggt 60
ggggtgctca tgggtggaggt gccttttccag ggaaggaccc taaccaaggt tgacagaagt 120
ggtgcctata tcgtgaggca ggctgcaaag agtggttggtg caaatggcct tgccagaaggt 180
tgcattgtcc aagtttcccta tgccattgggt gtccctgagc ccttgtcagt gtttgtggac 240
acttatggaa ctgggaagat tcctgacaag gagattcttc aaattgtgaa ggagaatttc 300
gacttcagac ctggaatgat c 321

<210> 1734

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1734

gtcgcnngca cgcgtacgta agctcggaa ttcggctcga ggacgagacc ccagaattga 60
tgccattgag tcatgttctt gcaactaaac tccgtgctcg tctcaccgag gttcgcaaga 120
acggaacctg cccatgggtg aggcctgatg ggaagaccca agtgactgtt gagtattaca 180
atgacaacgg tgccatgggt ccagttcgtg tccacactgt gcttatctcc acccaacatg 240
atgagactgt gaccaacgac gaaattgcag ctgacctcaa ggagcatgtg atcaagccgg 300
tgatcccgga 310

<210> 1735

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 1735

ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gtggatcatgg tctttggaga 60
gatcacaacc aaggccaacg tagactntga gaagattgtc cgtgacacat gccgcgaaat 120
tggaattcatc tctgatgatg ttggtcttga tgctgacaaa tgcaaggtgt tggtaacat 180
tgagcagcag agccctgata tcgcccaggg tgtgcacggt cacttcacca agcgcccaga 240
ggaggttggt gctggtgacc agggtcacat gtttggctat gccactga 288

<210> 1736
 <211> 299
 <212> nucleic acid
 <213> Glycine max

 <400> 1736

 gtcgcangca cgcgtacgtg agctcgggnt tcggctcgag ntcagtgaac gaggggcacc 60
 ctgacaagct ctgngaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc 120
 ctgacagcaa ggttgccctgt naaacctgca ccaagaccaa catggtgatg gttttcggag 180
 agatcacaac caaggccaac gtggactatg agaagattgt gcgtgacaca tgcaggaaca 240
 ttggttttgt ctctgatgat gttggtcttg atgctgacaa ctgcaaggtc ctcnncaan 299

<210> 1737
 <211> 328
 <212> nucleic acid
 <213> Glycine max

 <400> 1737

 ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggcctgtcat tcctgagaag 60
 taccttgatg agaagaccat cttccacctt aacccttctg gccgttttgt cattggtggc 120
 cctcatggtg atgctggtct cactggaaga aagatcatca ttgataccta tggtaggtgg 180
 ggtgctcatg gtggaggtgc cttttcaggg aaggacccta ccaaggttga cagaagtgg 240
 gcctatattg taaggcaggc tgcaaagagt gtcgtggcaa atggccttgc tagaagggtgc 300
 ttgtgcaagt ttccctatgc catggtgc 328

<210> 1738
 <211> 315
 <212> nucleic acid
 <213> Glycine max

 <400> 1738

 tcncgcgtac gtnagctcgg aattcggctc gagaccaaga ccaacatggg gatgggtttc 60
 ggagagatca caaccaaggc caacgtggac tatgagaaga ttgtgcgtga cacatgcagg 120
 aacattggtt ttgtctctga tgatgttggt cttgatgctg acaactgcaa ggtcctcgtc 180
 aacattgagc aacagagtcc tgatattgct caagggtgtgc acggccacct tcacaaagag 240

gcctgaggag attggtgctg gtgaccaagg tcatatgttc ggctatgccc actgacgaga 300
ctcccagact cagcc 315

<210> 1739
<211> 303
<212> nucleic acid
<213> Glycine max

<400> 1739

cccngcacgc gtacgtaagc tcggaattcg gctcgagcga tacttatgga ggatggggtg 60
ctcatgggtg tggtgctttc tcgggaagg accctaccaa ggttgatagg agtgggtgctt 120
acattgtgag acaggctgct aagagcattg tggcaagtgg acttgccaga aggtgcattg 180
tgcaagtgtc ttatgccatt ggtgtgcttg agcctttgtc tgtgtttgtt gacacctatg 240
gcactgggaa gatccatgat aaggagattc tcaacattgt gaaggaaaac tttgattcag 300
gcc 303

<210> 1740
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 1740

tctntgnanc gtagtaagct cggaattcgg ctcgagctga tattgcccag ggtgtgcacg 60
gccaccttac caaaagaccc gaggaatcg gtgctggaga ccagggtcac atgtttggct 120
atgccacgga cgagacccca gaattgatgc cattgagtca tgttcttgca actaaactcg 180
gtgctogtct caccgaggtt cgcaagaacg gaacctgccc atggttgagg cctgatggga 240
agacccaagt gactgttgag tattacaatg acaacggtgc catggttcca gttcgtgtc 299

<210> 1741
<211> 263
<212> nucleic acid
<213> Glycine max

<400> 1741

cattgagcaa cagagtcctg atattgctca aggtgtgcac ggccacctca caaagaggcc 60
tgaggagatt ggtgctggtg accaaggtca tatgttcggc tatgccactg acgagactcc 120

cgagctcatg cccttgagcc atgtccttgc cacgaagctc ggtgcccaagc tcaccgaggt 180
 tcggaagaac gggacatgcc cttggctgag acctgatggc aagacccaag tcaactgttga 240
 gtactacaat gacaaggggtg cca 263

<210> 1742
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 1742

gtngangcgt acgtaagctc ggaattcggc tcgaggcacg gccacctcac aaagaggcct 60
 gaggagattg gtgctgggtga ccaaggtcat atgttcggct atgccactga cgagactccc 120
 gagctcatgc ccttgagcca tgtccttgcc acgaagctcg gtgccaaagct caccgaggtt 180
 cggaagaacg ggacatgccc ttggctgaga cctgatggca agacccaagt cactgttgag 240
 tactacaatg acaaggggtgc catgggtcca atccgcgtcc acaactgtgt catctccac 299

<210> 1743
 <211> 254
 <212> nucleic acid
 <213> Glycine max
 <400> 1743

ctcaccgagg ttcgcaagaa cggtagctgc ccttggttga ggctgatgg gaagacccaa 60
 gtgaccgttg agtattacaa tgacaatggt gccagggttc ctattcgtgt acacaccgtg 120
 ctaatctcca cccaacacga cgagactgtc accaatgacg aaattgctgc tgacctcaaa 180
 gagcatgtga tcaagcctgt gatcccagag aagtaccttg atgagaagac cattttccac 240
 ttgaaccctt cagg 254

<210> 1744
 <211> 268
 <212> nucleic acid
 <213> Glycine max
 <400> 1744

acagagtcct gatattgctc aagggtgtgca cgccacctc acaaagaggc ctgaggagat 60
 tgggtgctggg gaccaaggtc atatgttcgg ctatgccact gacgagactc ccgagctcat 120

gcccttgagc catgtccttg ccacgaagct cggtgccaag ctcaccgagg ttcggaagaa 180
 cgggacatgc ccttggtgta gacctgatgg caagacccaa gtcactgttg agtactacaa 240
 tgacaagggt gccatggttc caatccgc 268

<210> 1745
 <211> 305
 <212> nucleic acid
 <213> Glycine max

<400> 1745

gcacgcgtac gtaagctcgg aattcggctc gagcacggac gagaccccag aattgatgcc 60
 attgagtcac gttcttgcaa ctaaactcgg tgctcgtctc accgaggttc gcaagaacgg 120
 aaactgcccc tgggtgaggg ctgatgggaa gacccaagtg actgttgagt attacaatga 180
 caacggtgcc atggttccag ttcgtgncca cactgtgctt atctccaccc aacatgatga 240
 gactgtgacc aacgacgaaa ttgcagctga cctcaaggag catgtgatca agccggtgat 300
 ccggg 305

<210> 1746
 <211> 316
 <212> nucleic acid
 <213> Glycine max

<400> 1746

antcgcangc acgcgtacgt aagctcggaa ttcggctcga ggtcctcgac gcttgccctg 60
 aacaggaccc agacagcaag gttgcctgcg aaacatgcac caagaccaac ttggtcatgg 120
 tcttcggaga gatcaccacc aaggccaacg ttgactacga gaagatcgtg cgtgacacct 180
 gcaggaacat cggcttcgtc tcaaacgatg tgggacttga tgctgacaac tgcaagggtcc 240
 ttgtaaacat tgagcagcag agccctgata ttgccagggt tgtgcacggc caccttacca 300
 aaagacccga ggaaat 316

<210> 1747
 <211> 306
 <212> nucleic acid
 <213> Glycine max

<400> 1747

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ctcaccggcc gcaagatcat 60
catcgacacc tatggaggat ggggtgcaca tgggtggtggt gccttctctg ggaaggatcc 120
taccaagggtt gataggagtgt gtgcctacat tgtgaggcaa gctgcaaaga gcattgttgc 180
aatggactt gctaggaggg caattgtgca agtttctat gccattggtg tgcctgagcc 240
cttgtctgtg tttgttgaca cttatggcac tgggaagatc cctgacaagg aaatcctcag 300
cattgt 306

<210> 1748
<211> 269
<212> nucleic acid
<213> Glycine max

<400> 1748

gtgcgtgaca catgcaggaa cattgggtttt gtctctgatg atgttgggtct tgatgctgac 60
aactgcaagg tcctcgtcaa cattgagcaa cagagtcctg atattgctca aggtgtgcac 120
ggccacctca caaagaggcc nnaggagatt ggtgctggtg accaagggtca tatgttcggc 180
tatgccactg acgagactcc cgagctcatg cccttgagcc atgtccttgc cacgaagctc 240
ggtgccaagc tcaccgaggt tcggaagaa 269

<210> 1749
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 1749

tcgcangcac ncgtacgtaa gctcggaatt cggctcgaga cagagtcctg atattgctca 60
aggtgtgcac ggccacctca caaagaggcc tgaggagatt ggtgctggtg accaagggtca 120
tatgttcggc tatgccactg acgagactcc cgagctcatg cccttgagcc atgtccttcc 180
acgaagctcg gtgccaagct caccgaggtt cggaagaacg ggacatgcc ttggctgaga 240
cctgatggca agaccaagt cactgttgag tactacaatg acaaggggtgc catggttcca 300
atccgcgtcc a 311

<210> 1750
<211> 308
<212> nucleic acid

<213> Glycine max

<400> 1750

gcangcacgc gtacgtaagc tcggaattcg gctcgangtt cttgcaacta aactcgggtgc 60
tcgtctcacc gaggttcgca agaacggaac ctgcccattg ttgaggcctg atgggaagac 120
ccaagtgact gttgagtatt acaatgacaa cgggtgccatg gttccagttc gtgtccacac 180
tgtgtttatc tccacccaac atgatgagac tgtgaccaac gacgaaattg cagtgcacct 240
aaggagcatg tgatcaagcc ggtgatcccg gagaagtacc ttgatgagaa gaccattttc 300
catttgaa 308

<210> 1751

<211> 394

<212> nucleic acid

<213> Glycine max

<400> 1751

aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 60
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 120
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 180
accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggtttttcg 240
gagagatcac aaccaaggcc aacgtggact atgagaagat tgtgcgtgac acatgcaagg 300
aacattggnt tttgtctctg atgaatgttg gncttgatgc tgacaactgc aaggnccccc 360
tcaaanattg gnnacaaaa ntccggaana ttgc 394

<210> 1752

<211> 326

<212> nucleic acid

<213> Glycine max

<400> 1752

cangcacgcg tacgtaagct cggaattcgg ctcgagggac cctaccaagg ttgacagaag 60
tggtgcctat atcgtgaggc aggctgcaaa gagtgttgtg gcaaatggcc ttgccagaag 120
gtgcattgtc caagtttccct atgccattgg tgtccctgag cccttgtcag tgtttgtgga 180
catttatgga actgggaaga ttccctgacaa ggagattcctt caaattgtga aggagaattt 240

cgacttcaga cctggaatga tcaccattaa cttggacctt aagaggggtg gccatagggt 300
cctcaagaca gctgcttatg gacact 326

<210> 1753
<211> 536
<212> nucleic acid
<213> Glycine max

<400> 1753

gnnggngngt gnnttcntnn nnnntnacnn tttggcntgc cgtaccggtc cggaattccc 60
gggtcgaccc acgcgtccgg caagccccac tcaaccacca cacctcttct cgttcacgct 120
acccctttct gctcttcttc tacctttcaa gttttaaaag tataaagatg gcagagacat 180
tcctatttac ctcagagtcg gtgaacgagg gacacctga caagctctgc gaccaaactc 240
ccgatgctgt cctcgacgct tgccctcgagc aggaccaga cagcaaagt gcctgcgaaa 300
catgcaccaa aaccaacttg gtcattggtc tcggagaaat cagcaccaag gccaacgttg 360
actacgagaa gatagtgcgt gacacctgca ggaacatcgg cttcgtctca aatgatgtgg 420
gactggatgc cgacaactgc aagggtctgt caacattgac agcagaccct gatattggtc 480
aagggtggtc acgggcacct taccaaaaaa anctggaaga aattgggggtc tggnga 536

<210> 1754
<211> 286
<212> nucleic acid
<213> Glycine max

<400> 1754

cacgcgtacg taagctcgga attcggctcg agaccaaggt tgataggagt ggtgcttaca 60
ttgtgagaca ggctgctaag agcattgtgg caagtggact agccagaagg tgcattgtgc 120
aagtgtctta tgccattggt gtgcccagac ctttgtctgt ctttgttgac acctatggca 180
ccgggaagat ccatgataag gagattctca acattgtgaa ggagaacttt gatttcaggc 240
ccggtatgat ctccatcaac cttgntctca agaggggtgg gaataa 286

<210> 1755
<211> 276
<212> nucleic acid
<213> Glycine max

<400> 1755

tagtgcctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg gactatgaga agattgtgcg 240
tgacacatgc aggaacattg gttttgtctc tgatga 276

<210> 1756

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 1756

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag ggccaacgta gactatgaga 60
agattgtccg tgacacatgc cgcgaaattg gattcatctc tgatgatgtt ggtcttgatg 120
ctgacaaatg caaggtgttg gtcaacattg agcagcagag ccctgatatc gcccagggcg 180
tgacagggtc attcaccaag cgcacagagg aggttggtgc tggtgaccag ggtcacatgt 240
ttgggctatg ccatgatgaa acccctgagt acatgcccct cagccatgtc cttgcaacca 300

<210> 1757

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 1757

nngtcgcang cacgcgtacg taagctcgga attcggctcg agccctgata tcgcccaggg 60
tgtgcacggg cacttcacca agcgcacaga ggaggttggt gctggtgacc agggtcacat 120
gtttggctat gccactgatg aaaccctga gtacatgcc ctcagccatg tccttgcaac 180
caaactcggg gctgcctca ccgaggttag gaaaaatggg acctgtgctt ggctgaggcc 240
agatggcaag acacaagtaa ctgttgagta ctacaatgac aatggtgcca tggttccagt 300
tcgt 304

<210> 1758

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 1758

ngtcgcangc acgcgtacgt aagctcggaa ttcggctcga gctgcaaaga gcattgttgc 60
aaatggactt gctaggaggg caattgngca agtttcctat gccattggtg tgctgagcc 120
cttgtctgtg tttgttgaca cttatgggca ctgggaagat ccctgacaag gaaatcctca 180
gcattgtgaa ggagagtttt gacttcaggc ctggcatgat ctccatcaac cttgatctca 240
agaggggtgg aaatggcagg ttcttgaaga ctgctgcata tggacacttt ggagagatg 300
accctgact 309

<210> 1759

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 1759

ncacgtcgan gcacgcgtac gtaagctcgg aattcggtc gaggagaaga ccatcttcca 60
ccttaaccct tctggccgtt ttgtcattgg tggccctcat ggtgatgctg gtctcactgg 120
aagaaagatc atcattgata cctatggtgg gtgggggtgct catggtggag gtgccttttc 180
aggggaaggac cctaccaagg ttgacagaag tgggtgcctat atcgtgagggc aggctgcaaa 240
gagtgtttgt gcaaattggcc ttgccagaag gtgcatgtcc aagtttccta tgccattggt 300
gtccctgagc cctgtcagtg 320

<210> 1760

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 1760

gtcgcangca cgcgtacgta agctcggaa tccgctcgag gcgtgacaca tgcaggaaca 60
ttggttttgt ctctgatgat gttggtcttg atgctgacaa ctgcaaggctc ctctgcaaca 120
ttgagcaaca gagtctgat attgctcaag gtgtgcacgg ccacctcaca aagaggcctg 180
aggagattgg tgctggtgac caaggtcata tgttcggcta tgccactgac gagactcccg 240
agctcatgcc cttgagccat gtccttgcca cgaagctcgg tgccaagctc accgn 295

<210> 1761
 <211> 297
 <212> nucleic acid
 <213> Glycine max

 <400> 1761

 ngtcntangc acgcgtacgt aagctcggaa ttcggctcga gggtaacat tgagcaacag 60
 agcccggata tcgccaggg tgtgcacggc cacttcacca agcggccaga ggaggttggt 120
 gctgggtgacc agggtcacat gtttgggtat gccaccgatg aaacccccga gtacatgccc 180
 ctcagccatg tccttgcaac caaacttggt gctcgctca cagaggtag gaagaatggc 240
 acctgtgctt ggttgaggcc agatggtaag acacaagtaa ccgtcgagta ctacaat 297

<210> 1762
 <211> 297
 <212> nucleic acid
 <213> Glycine max

 <400> 1762

 tcgcatncac gcgtacgtaa gctcggaatt cggctcgagc atcattgata cctatggtgg 60
 ctgggggtgct catggtggag gtgccttttc aggggaaggac cctaccaagg ttgacagaag 120
 tgggtgcctat attgtaaggc aggctgcaaa gagtgctctg gcaaattggcc ttgctagaag 180
 gtgcattgtg caagtttctt atgccattgg tgtccctgag cccttgtcag tgtttgtgga 240
 cacttatgga actgggaaga ttcttgacaa ggagattctg caaattgtga aggagaa 297

<210> 1763
 <211> 303
 <212> nucleic acid
 <213> Glycine max

 <400> 1763

 angangacgt cgctgcacgc gtacgtaagc tcggaattcg gctcgaggga agaacgggac 60
 atgcccttgg ctgagacctg atggcaagac ccaagtcact gttgagtact acaatgacaa 120
 gggtgccatg gttccaatcc gcgtccacac tgtgctcatc tccacacagc atgatgagnc 180
 tgtcacaaat gatgagattg cagctgatct taaagaacac gtgattaagc ctgtgattcc 240
 tgagaagtac cttgatgaga agaccatttt ccatttgaac ctttctggca ggtttgtcat 300

tgg

303

<210> 1764
<211> 492
<212> nucleic acid
<213> Glycine max

<400> 1764

gggnaantct acgcgnnacg ttctggtcag agccattccc gggananaacc cacgcnccn 60
tacggctgcg agaaagacga cagaaggggg caacgctttg attttgaggg caaggcaaag 120
ccccactcaa accaacacac ctctcctccg ttcacgctac cttttctgct cttctttctac 180
ctttcaagtt ttaaaaagta taaagatggc agagacattc ctattttacct cagagtcggt 240
gaacgagggga caccctgaca agctctgcca ccaaattctc gatgctgtcc tcgacgcttg 300
cctcgagcag gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt 360
catggtcttc ggagaaatca cgaccaaggc caacgttgac tacgagaaga tagtgcggtga 420
cacctgcagg aacatcggtc tcgtctcaaa tgatgtggga ctggatgcgc acaactgcaa 480
aggtcctcgt ca 492

<210> 1765
<211> 295
<212> nucleic acid
<213> Glycine max

<400> 1765

acgtcgcang cacgctgacg taagctcgga attcggctcg aggtgccttt tcagggaagg 60
accctaccaa gggtgacaga agtggtgcct atatcgtgag gcaggctgca aagagtgttg 120
tggcaaattg ccttgccaga aggtgcattg tccaagtttc ctatgccatt ggtgtccctg 180
agcccttgtc agtgttttgtg gacacttatg gaactgggaa gattcctgac aaggagattc 240
ttcaaattgt gaaggagaat ttcgacttca gacctggaat gatcaccatt aactt 295

<210> 1766
<211> 290
<212> nucleic acid
<213> Glycine max

<400> 1766

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gaagctctgt gaccagatct 60
ccgatgctgt gctcgatgca tgcttggagc aggaccctga cagcaagggt gcctgtgaaa 120
cctgcaccaa gaccaacatg gtgatgggtt tcggagagat cacaaccaag gccaacgtgg 180
actatgagaa gattgtgcgt gacacatgca ggaacattgg tttgtctct gatgatgta 240
ntcttgatgc tgacaactgc aaggctcctg tcaacattga gcaacagagt 290

<210> 1767
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 1767

620

nagtcnncng aacgcgtngg taagctcggg anttcggctc gangtgctca tgggtggangn 60
gccttttcag ggaaggaccc taccaagggt gacagaagtg gtgcctatat cgtgaggcag 120
gctgcaaaga gtgttgtggc aaatggcctt gccagaagggt gcattgtcca agtttcctat 180
gccattgggtg tccttgagcc cttgtcagtg tttgtggaca cttatggaac tgggaagatt 240
cctgacaagg agattcttca aattgtgaag gagaatttcg acttcagacc tggaatgatc 300

<210> 1768
<211> 327
<212> nucleic acid
<213> Glycine max
<400> 1768

gcagtgtacg tnagctcggg atttcggctc agcattgggt tgcctgagcc cttgtctgtg 60
tttgttgaca cttatggcac tgggaagatc cctgacaagg aaatcctcag cattgtgaag 120
gagagttttg acttcaggcc tggcatgatc tccatcaacc ttgatctcaa gaggggtgga 180
aatggcagggt tcttgaagac tgcngcatat ggacactttg gcagagatga ccctgacttc 240
acatgggaag tggatgaagc actcaagggg gaaaaggtag tgcttaacta aaaggggttc 300
caacactctt ggccaangga ttttgcc 327

<210> 1769
<211> 322
<212> nucleic acid
<213> Glycine max

<400> 1769

gcgtacgnaa gctcgggaatt cggctcgaga atggacttgc taggagggca attgtgcagt 60
ttcctatgcc attggtgtgc ctgagccctt gtctgtgttt gttgacactt atggcactgg 120
gaagatccct gacaaggaaa tcctcagcat tgtgaaggag agttttgact tcaggcctgg 180
catgatctcc atnaaccttg atctcaagag ggggtggaaat ggcaggttct tgaagactgc 240
tgcatatgga cactttggca gagatgacct tgacttcaca tgggaagtgg tgaagccatc 300
aagggggaga agacctgctt aa 322

<210> 1770

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 1770

gnncgnncgn aagctcggaa ttcggtcga gngccaagct caccgagggt cggaagaacg 60
ggacatgcc ttggctgaga cctgatggca agaccaagt cactgttgag tactacaatg 120
acaaggggtgc catggttcca atccgcgtcc aactgtgtct catctccaca cagcatgang 180
agnctgtcac aaatgatgag attgcagctg atcttaaaga acacgnnatt aagcctgtna 240
tncnganaa gtnccttnat gagaagacca ttttccattt gaacccttc 289

<210> 1771

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 1771

nnngcntnan gtagcgnacg taagctcgga attcggtcgc aggaagaac gggacatgcc 60
cttggctgag acctgatggc aagaccaag tcaactgttg gtactacaat gacaaggggtg 120
ccatggttcc aatccgcgtc cacactgtgc tcatctccac acagcatgat gagactgtca 180
caaatgatga gattgcagct gatcttaaag aacacgtgat taagcctgtg attcctgaga 240
agtaccttga tgagaagacc attttccatt tgaacccttc tggcagggtt gtcattg 297

<210> 1772

<211> 260

<212> nucleic acid

<213> Glycine max

<400> 1772

catgcaccaa gaccaacatg gtcatgggtct ttggagagat cnntaccaag gccaacgtag 60
actatgagaa gattgtccgt gacacatgcc gcgaaattgg attcatctct gatgatgttg 120
gtcttgatgc tgacaatgca aggtgtttgg caacattgag cagcagagcc ctgatatcgc 180
ccaggggtgtg cacgggtcact tcaccaagcg ccagaggag gttggtgctg gtgaccaggg 240
tcacatgttt ggctatgcca 260

<210> 1773

<211> 338

<212> nucleic acid

<213> Glycine max

<400> 1773

ccnnccccca ccnncntacn aaagctcgga attcggtctg aggttcgggt atgccactga 60
cgagactccc gagctcatgc ncttgagcca tgtccttgcc acgaagctnc ggtgccaaagc 120
tcaccgaggt tcggaanaac gggacatgcc cttggctgag acctgatggc aagaccaag 180
tcactgttga gtactacaat gacaaggggt ccatggttcc aatccgcgtc cacactgtgc 240
tcctctccac acagcatgat gagactgtca caaatgatga gattgcagtg atcttaaaga 300
acacgtgatt aagcctgtga ttnttgagaa gtaccttg 338

<210> 1774

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 1774

tngnangcac gcgtacgnaa gctcgggaatt cggctcgaga actttcctat tcacatctga 60
atcagtgaac gaggggcacc ctgacaagct cctgtgacca gatctccgat gctgtgctcg 120
atgcatgctt ggagcaggac cctgacagca aggttgcttg tgaaacctgc accaagacca 180
acatggtgat ggttnccgga gagatcacia ccaaggccaa cgtggactat gagaagattg 240
tgcgtgacac atgcaggaa attggttttg tctctgatga tgttggtctt gatg 294

<210> 1775

<211> 317
 <212> nucleic acid
 <213> Glycine max

 <400> 1775

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 gacaagctgt gcgaccagat ctctgatgca gtgctcgatg ngtgcnttga acaggaccct 120
 gacagcaagg ttgcntgtga gacatgcacc aagaccaaca tggatcatggt ctttggagag 180
 atcacaacca aggccaacgt agactatgag aagattgtcc gtgacacatg ccgcgaaatt 240
 gggattcatc tctgggtggtg ttggtcttga tncgtgacaat gcaaggtgnt ggtcaaacat 300
 tgagcagcag agccctg 317

<210> 1776
 <211> 309
 <212> nucleic acid
 <213> Glycine max

 <400> 1776

 gtgcgacgca cgcgtacgta agctcgggaat tcggctcgag gaccattttc catttgaacc 60
 cttctggcag gtttgtcagc ggagggccgc atggcgatgc tggctctacc ggccgcaaga 120
 tcatcatcga cacctatgga ggatggggtg cacatgggtg tggtgccctc tctgggaagg 180
 atcctaccaa ggttgatagg agtgggtgcct acattgtgag gcaagctgca aagagcattg 240
 ttgcaaattg acttgctagg aggcaattgt gcaagtttcc tatgccattg gtgtgcctga 300
 gcccttgtc 309

<210> 1777
 <211> 329
 <212> nucleic acid
 <213> Glycine max

 <400> 1777

 gtgcgcatgca cgcgtacgta agctcgggaat tcggctcgag gcagggaccc agacagcaaa 60
 gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg tcttncggag aaatcacgac 120
 caaggccaac gttgactacg agaagatagt gcgtgacacc tgcaggaaca tcggcttcgt 180
 ctcaaatgat gtgggactgg atgccgacaa ctgcaaggtc ctogtcaaca ttgagcagca 240

gagccctgat attgctcagg gtgtacacgg ccaccttacc aaaaaacctg aagaaattgg 300
 tgctggtgac cagggtcaca tgtttggct 329

<210> 1778
 <211> 518
 <212> nucleic acid
 <213> Glycine max
 <400> 1778

gaggnnnnna gggntnntnn tatgaancna nggaactttt nngcntgccc gtaccgggtcc 60
 ggattccccg gtcgacccac gcgtccgtac ggctgcggaa gacgacagaa gggggcagcg 120
 cttgatttga ggccaggcaa gcccactca accaccacac ctctcctcgt tcacgtacc 180
 cttttctgct cttcttctac ctttcaagtt ttaaaagtat aaagatggca gagacattcc 240
 tattttacctc agagtcggtg aacgagggac accctgacaa gctctgcgac caaatctccg 300
 atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc tgcgaaacat 360
 gcacaaaaac caacttggtc atggtcttcg gagaaatcac gaccaaggcc aacgttgact 420
 acgagaagat agtgcgtgac acctgcagga acatcggett cgtctcaaata gatgtgggac 480
 tggatgcccg acaactgcaa ggtcctcgtt acattgac 518

<210> 1779
 <211> 293
 <212> nucleic acid
 <213> Glycine max
 <400> 1779

gtcgcangca cgcgtacgta agctcggaat tcnctcgag ntgagcaatg accaaattgc 60
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 gaccatcttc caccttaacc cttctggccg ttttgtcatt ggtggccctc atggtgatgc 180
 tgggtctcact ggaagaaaga tcatcattga tacctatggt ggggtggggtg ctcatggtgg 240
 aggtgccttt tcaggggaagg accctaccaa ggttgacaga agtgggtgcct ata 293

<210> 1780
 <211> 269
 <212> nucleic acid
 <213> Glycine max

<400> 1780

cgagggacac cctgataagc tctgcgacca aatctccgat gctgtcctcg acgcttgcct 60
cgaacaggac ccagacagca aggttgctcg cgaaacatgc accaagacca acttggtcat 120
ggtcttcgga gagatcacca ccaaggccaa cgttgcatac gagaagatcg tgcgtgacac 180
ctgcaggagc atcggcttca tctcanacga tgtgggactt gatgctgaca actgcaaggt 240
ccttgtnaac attgagcagc ngagccctg 269

<210> 1781

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 1781

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagg tgggggtgctc atggtggagg 60
tgccctttca gggaaggacc ctaccaaggt tgacagaagt ggtgcctata tcgtgaggca 120
ggctgcaaag agtggttggt caaatggcct tgccagaagg tgcattgtcc aagtttccta 180
tgccattggt gtccctgagc ccttgctcagt gtttgtggac acttatggaa ctgggaagat 240
tcctgacaag gagattcttc aaattgtgaa ggagaatttc gacttca 287

<210> 1782

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 1782

cgngcacgc gtacgtnagc tcggaattcg gctcgaggca cggccacctc acaaagaggc 60
ctgaggagat tgggtgctggt gaccaaggct atatgttcgg ctatgccact gacgagactc 120
ccgagctcat gcccttgagc catgtccttg ccacgaagct cggtgccaag ctcaccgngg 180
ttcggaanaa cgggacatgn ccttggtgta nacctgatgg caagacncaa gtcactgttg 240
agtactacaa tgacaagggt gccatgggtc caatccgctg ccacactgtg ctcactctcca 300
c 301

<210> 1783

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 1783

nagtcgcang cacgcgtacg taagctcgga attcggctcg agccagaatt catgccattg 60
agtcattgttc ttgcaaccaa gctcgggtgct cgtctcaccg aggttcgcaa gaacggaacc 120
tgcccatggc tgaggcctga tgggaagacc caagtgactg tggagtatta caatgataat 180
ggtgccaggg ttccagttcg tgtncacacc gtgctaattc ccaccagca tgatgagact 240
gtcaccaacg acgaaattgc ggctgacctc aaggagcatg tgatcaagcc tgtgatcccg 300
gagaa 305

<210> 1784

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 1784

gtcgtgcac gcgtacgtna gctcggaatt cggctcgagt gatgatgttg gtcttgatgc 60
tgacaaatgc aagggtgttg tcaacattga gcagcagagc cctgatatcg ccagggtgt 120
gcacggtcac ttcaccaagc gccagagga ggttggtgct ggtgaccagg gtcacatggt 180
tggctatgcc actgatgaaa ccctgagta catgcccctc agccatgtcc ttgcaaccaa 240
actcgggtgct cgcctcaccg aggttaggaa aaatgggtacc tgtgctt 287

<210> 1785

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1785

nancacgtcg cangcacgcg tacgtaagct cgggaattcg gctcgagagg gtcacatggt 60
tggctatgcc actgatgaaa ccctgagta catgcccctc agccatgtcc ttgcaaccaa 120
actcgggtgct cgcctcaccg aggttaggaa aaatgggtacc tgtgcttggc tgaggccaga 180
tggcaagaca caagtaactg ttgagtacta caatgacaat ggtgcatgg ttccagttcg 240
tgtccacact gtcctaattt ccacccaaca tgatgagnct gtgagcaatn accaaattgc 300
tgctgacctt 310

<210> 1786
 <211> 287
 <212> nucleic acid
 <213> Glycine max

 <400> 1786

 gtgcgacgca cgcgtacgta agtcggaat tcggctcgag ntcacaacca aggccaacgt 60
 agactatgaa aagattgtcc gcgacacatg ccgcgaaatt ggattcatct ctgatgatgt 120
 tgggtcttgat gctgacaaat gcaaggtgtt ggtcaacatt gagcaacaga gcccggatat 180
 cggccagggt gtgcacggcc acttcaccaa gcgcccagag gaggttggtg ctggtgacca 240
 gggtcacatg tttgggtatg ccaccgatga aacccccgag tacatgc 287

<210> 1787
 <211> 295
 <212> nucleic acid
 <213> Glycine max

 <400> 1787

 gtgcgatgca cgcgtacgta agtcggaat tcggctcgag gtcttgatgc tgacaactgc 60
 aaggctctcg tcaacattga gcaacagagt cctgatattg ctcaaggtgt gcacggccac 120
 ctcacaaaga ggcttgagga gattggtgct ggtgaccaag gtcatatgtt cggctatgcc 180
 actgacgaga ctcccgagct catgcccttg agccatgtcc ttgccacgaa gctcggtgcc 240
 aagctcaccg aggttcggaa gaacgggaca tgcccttggc tgagacctga tggca 295

<210> 1788
 <211> 321
 <212> nucleic acid
 <213> Glycine max

 <400> 1788

 tcgcacgcac gcgtacgtaa gctcgggaatt cggctcgagn agatggcaga gacattccta 60
 tttacctcag antcgggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat 120
 gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 180
 accaaaacca acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 240
 gagaagatag tgcgtgacac ctgcaggaac atcggcttcg tctcaaatga ngtgggactg 300

gatgccgaca actgcaagtc t

321

<210> 1789

<211> 270

<212> nucleic acid

<213> Glycine max

<400> 1789

tagtgccctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60

agtgaacgag gggcacccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120

atgcttgag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180

ggtgatggtt ttcggagaga tcacaaccaa ggccaacgtg gactatgaga agattgtgcg 240

tgacacatgc aggaacattg gttttgtctc 270

<210> 1790

<211> 333

<212> nucleic acid

<213> Glycine max

<400> 1790

tngatanacg cgtacgtgag ctccgaattc ggctcgaggg aggttgggtgc tgggtgaccag 60

ggtcacatgt ttgggctatg ccactgatga aaccctgag tacatgcccc tcagccatgt 120

cttgcaacca aactcgggtgc tcnctcacc gaggttagga aaaatggtac ctgtgcttgg 180

ctgaggccag atggcaagac acaagtaact gttgagtact acaatgacaa tgggtgccatg 240

gttccagttc gtgtccacat gtcctaattt ccaccaacaa tgatgagcct gtgagcaatg 300

accaaattgc tgctgacctt aaagagcatg tta 333

<210> 1791

<211> 267

<212> nucleic acid

<213> Glycine max

<400> 1791

caatggtgcc atggttccag ttcgtgtcca cactgtccta atttccaccc aacatgatga 60

nacctgtgag caatgaccaa attgctgctg acctaaaga gcatgttatc aagcctgtca 120

ttcttgagaa gtacctggat gagaagacca tcttccacct taacccttct ggccgttttg 180

tcattggtgg ccctcatggt gatgctggtc tcaactggaag aaagatcatc attgatacct 240
atggtgggtg ggggtgctcat ggtggag 267

<210> 1792
<211> 314
<212> nucleic acid
<213> Glycine max
<400> 1792

ccanaatcgc atgcacgcgt acgtaagctc ggaattcngc tcgagctcga gccgctcgag 60
ccggaatcag tgaacgaggg gcaccctgac aagctctgtg accagatcct ccgatgctgt 120
gctcgatgca tgcttgagc aggaccctga cagcaagggt gctgtgaaa cctgcaccaa 180
gaccaacatg gtgatggttt tcggagagat cacaaccaag gccaacgtgg actatgagaa 240
gattgtgcgt gacacatgca ggaacattgg ttttgtctct gatgatgttn gtcttgatgc 300
tgacaactgc aagt 314

<210> 1793
<211> 512
<212> nucleic acid
<213> Glycine max
<400> 1793

gnnnnnaatt ctacgccggn ctctaacgcg nnaacanaat tcccggaaac gacccacgng 60
nccnntgcga gaagacgaca gaagggggca acgcttnagc agacttnaca acancacaaa 120
gcnngttact gtctgttcaa gctaacatct ccctctctct ttccttaant gcctccttnc 180
caagaaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcaa gttaacgaag 240
gggcaccccc gacaagctct gtgaccaaga tctccgatgc tgtgctcgat gcatgcttgg 300
agcaagacct tgacagcaan gttgcctgtg aaacctgcac caagaccaac atggtgatgg 360
ttttcggaga gattacaacc aangccaacg tggactatga gaagattgtg cgttacacat 420
gcangaacat tggttttgtc tctgaagatg ttggctctga agctgacaac tgcaangtcc 480
tcgtcaacaa ttaacaacaa naattctgat at 512

<210> 1794
<211> 294
<212> nucleic acid

<213> Glycine max

<400> 1794

nnngntctan gcacgcgtac gtaagctcgg aattcggctc gaggaccatc ttccacctta 60
accottcttg ccgttttctc attggtggcc ctcatggtga tgctggtctc actggaagaa 120
agatcatcat tgatacctat ggtggctggg gtgctcatgg tggagggtgcc ttttcaggga 180
aggaccctac caagggtgac agaagtggtg cctatatgtt aaggcaggct gcaaagagtg 240
tcgtggcaaa tggccttgct agaaggtgca ttgtgcaagt ttcctatgcc attg 294

<210> 1795

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 1795

cgannacgtc gcangcacgc gtacgtaagc tcggaattcg gctcgaggga cgagacccca 60
gaattgatgc cattgagtca tgttcttgca actaaactcg gtgctcgtct caccgagggt 120
cgcaagaacg gaacctgccc atggttgagg cctgatggga agaccaagt gactgttgag 180
tattacaatg acaacggtgc catggttcca gttcgtgtcc aactgtgtct tatctccacc 240
caacatgatg agactgtgac caacgacgaa attgcagctg acctcaagga gcatgtgatc 300
a 301

<210> 1796

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 1796

gcatgcacgc gtacgtaagc tcggaattcg gctcgagagt ggtgcctaca ttgtgaggca 60
agctgcaaag agcattgttg caaatggact tgctaggagg gcaattgtgc aagtttccta 120
tgccattggt gtgcctgagc ccttgtctgt gtttgttgac acttatggca ctgggaagat 180
ccctgacaag gaaatcctca gcattgtgaa ggagagtttt gacttcaggc ctggcatgat 240
ctccatcaac cttgatctca agaggggtgg aaatggc 277

<210> 1797

<211> 297
 <212> nucleic acid
 <213> Glycine max
 <400> 1797
 gcangcacgc nnacgtnagc tcggaattcg gctcgagacc ctaccaaggt tgacagaagt 60
 ggngcctata tcgtgaggca ggtgcaaaag agtggtgtgg canatggcct tgccagaagg 120
 tgcattgtcc aagtttccta tgccattggt gtccctgagc ccttgtcagt gtttgtggac 180
 acttatggaa ctgggaagat tcctgncaag gagattcttc aaattgtgaa ggagaatttc 240
 gacttnagac ctggaatgat caccattaac ttggacctta agaggggtgg ccatagg 297

<210> 1798
 <211> 264
 <212> nucleic acid
 <213> Glycine max
 <400> 1798
 agagcccgga tatcgcccag ggtgtgcacg gccacttcac caagcgccca gaggagggtg 60
 gtgctggtga ccagggtcac atgtttgggt atgncaccga tgaaaccccc gagtacatgc 120
 ccctcagcca tgtccttgca accaaacttg gtgctcgcct cacagagggt aggaagaatg 180
 gcacctgtgc ttggttgagg ccagatggta agacacaagt aaccgtcgag tactacaatg 240
 acaatggtgc catggttcca ttgc 264

<210> 1799
 <211> 311
 <212> nucleic acid
 <213> Glycine max
 <400> 1799
 nagtcgcang cacgcgtacg taagctcgga attcggctcg agccagaatt catgccattg 60
 agtcatgttc ttgcaaccaa gctcgggtgct cgtctcaccg aggttcgcaa gaacggaacc 120
 tgcccatggc tgaggcctga tgggaagacc caagtgactg tggagtatta caatgataat 180
 ggtgccaggg ttccagttcg tgnacacacc gtgctaactc ccaccagca tgatgagact 240
 gtcaccaacg acgaaattgc ggctgacctc aaggagcatg tgatnaagcc tgtgatcccg 300
 gngaagtnct t 311

<210> 1800
 <211> 508
 <212> nucleic acid
 <213> Glycine max

 <400> 1800

 ttttgtntnc cgaggcccn acnancntcn tggcctcgng nccacgcgta ngtaaantcg 60
 gaattcggct cgagatttga ggccaggcaa gcccactca accaccacac ctctcctcgt 120
 tcacgctacc cttttctgct cttctctac ctttcaagtt ttaaaagtat aaagatggca 180
 gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa gctctgcgac 240
 caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc 300
 tgcgaaacat gcacaaaaac caacttggtc atggtcttcg gagaaatcac gaccaaggcc 360
 aacgttgact acganaagat agtgcgtgac anctgcaaga acatcggctt cntctcaaatt 420
 gatgtgggac tggatgccga caactgcaag gtcctccgtc aacantgaac aacaagaacc 480
 ctgatattgc ncaagggttt naaccggc 508

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<210> 1801
 <211> 292
 <212> nucleic acid
 <213> Glycine max

 <400> 1801

 gtgcgancga cgcgtacgta agctcggaat tcggctcgag ccctaccaag gttgacagaa 60
 gtggtgacta tatcgtgagg caggctgcaa agagtgttgt ggcaaatggc cttgccagaa 120
 ggtgcattgt ccaagtttcc tatgccattg gtgtccctga gcccttgta gtgtttgtgg 180
 acacttatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 240
 togacttcag acctggaatg atcaccatta acttggaact taagaggggt gg 292

<210> 1802
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 1802

 cgtcgcangc acgcgtacgt aagctcgga ttcggctcga gctcgagccg cgcccagagg 60

aggttggtgc tggtgaccag ggtcacatgt ttggctatgc cactgatgaa acccctgagt 120
 acatgcccct cagccatgtc cttgcaacca aactcgggtgc tcgcctcacc gaggttagga 180
 aaaatggtac ctgtgcttgg ctgaggccag atggcaagac acaagtaact gttgagtact 240
 acaatgacaa tggtgccatg gttccagttc gtgtccacac tgtcctaatt tccaccaaac 300
 atgatac 306

<210> 1803
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 1803

acgtcgcattg cagcgtacg taagctcgga attcggctcg agctccgatg ctgtgctcga 60
 tgcattgcttg gagcaggacc ctgacagcaa ggttgcttgt gaaacctggc accaagacca 120
 acatggtgat ggttttcgga gagatcacia ccaaggccaa cgtggactat gagaagattg 180
 tgcgtgacac atgcaggaac attggtttta nctctgatga tgttggtctt gatgctgaca 240
 actgcaaggc cctcgtcaac attgagcaac agagtcctga tattgctcaa ggtgtgcacg 300
 gccacctca 309

<210> 1804
 <211> 437
 <212> nucleic acid
 <213> Glycine max
 <400> 1804

gagaagacga cagaaggggg cancgcttga ttnaggcca ggcangcccc actcanccac 60
 cacacctctc ctenttcacg ctaccctttt ctgctcttct tctanctttc aagttttaaa 120
 agtataaaga tggcagagan attcntatth acctcagagt cggatgaacga gggacaccct 180
 gacaagctct ggcaccaaatt ctccgatgct gtccctcgacg cttgcctcna gcaggancca 240
 nacagcaaaa ttgcctgcna aacatgcacc aaaaccaact tggatcatggt cttcggagan 300
 atcacgacca aggccaacgt tgactacgag aagatagtgc gtgacacctg caggaacatc 360
 ggcttcgtct caaaatgatg tgggactgga tcccgacaac tgcaangtcc tcgtcaacat 420
 ttgaacanca naggcct 437

<210> 1805
 <211> 299
 <212> nucleic acid
 <213> Glycine max

 <400> 1805

 gtcgcganna cgcgtacgta agctcggaat tcggctcgag aacagagtcc tgatattgct 60
 caaggtgtgc acggccacct tcacaaagag gcctgaggag attggtgctg gtgaccaagg 120
 tcatatgttc ggctatgcca ctgacgagac tcccgagctc atgcccttga gccatgtcct 180
 tgccacgaag ctcggtgcca agctcaccga ggctcggaag aacgggacat gcccttggct 240
 gagacctgat ggcaagaccc aagtcactgt tgagtactac aatgacaagg gtgccatgg 299

<210> 1806
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 1806

 cgtcgcatgc acgcgtacgt aagctcgga ttcggctcga gtcacgacca aggccaacgt 60
 tgactacgag aagatagtgc gtgacacctg caggaacatc ggcttcgtct caaatgatgt 120
 gggactggat gccgacaact gcaaggctct cgtcaacatt gagcagcaga gccctgatat 180
 tgctcagggt gtacacggcc accttaccaa aaaacctgaa gaaattgggtg ctggtgacca 240
 gggtcacatg tttggctatg cactgatga aaccttgaa ttgatgccat tgagcc 296

<210> 1807
 <211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 1807

 ntgcgatgca cgcgtacgta agctcggaat tcggctcgag ctcgagccgc attcctgaga 60
 agtacottga tgagaagacc atcttccacc ttaacccttc angccgtttt gtcattgggtg 120
 gccctcatgg tgatgctggt ctactggaa gaaagatcat cattgatacc tatggtggct 180
 ggggtgctca tgggtggagg gccttttcag ggaaggaccc taccaagggt gacagaagtg 240
 gtgcctatat tgtaaggcag gctgcaaaga gtgtcgtggc aaatggcctt gctagaagggt 300

gcatgtgc

308

<210> 1808
<211> 261
<212> nucleic acid
<213> Glycine max
<400> 1808

caggctgcta agagcattgt ggcaagtgga cttgccagaa ggtgcattgt gcaagtgtct 60
tatgccattg gtgtgcctga gcctttgtct gtgtttgttg acacctatgg cactgggaag 120
atccatgata aggagattct caacattgtg aaggaaaact ttgatttcag gcctgggtatg 180
atctccatca accttgatct caagaggggt ggaaataacn ggttttggan nactgccncc 240
natggacant tggaangnac c 261

<210> 1809
<211> 275
<212> nucleic acid
<213> Glycine max
<400> 1809

cgtcgcangc acgcgtacgt aagctcgga ttcggctcgn gctgagccct tgtctgtgtt 60
tgttgacact tatggcaactg ggaagatccc tgacaaggaa atcctcagca ttgtgaagga 120
gagttttgac ttcaggcctg gcatgatctc catcaacctt gatctcaaga ggggtggaaa 180
tggcagggttc ttgaagactg ctgcatatgg acactttggc agagatgacc ctgacttcac 240
atgggaagtg gtgaagccac tcaaggggga gaagg 275

<210> 1810
<211> 270
<212> nucleic acid
<213> Glycine max
<400> 1810

nantcgnang ctcngaattc ggctcgaggg tgatggtttt cggagnnntc anaaccaagg 60
ccaacgtgga ctatgagaag attgtgcgtg acacatgcag gaacattggt tttgtctctg 120
atgatgttgg tcttgatgct gacaactgca aggtcctcgt caacattgag caacagagtc 180
ctgatattgc tcaaggtgtg cacggccacc tcacaaagag gcctgaggag attggtgctg 240

gtgaccaagg tcatatgttc ggctatgcca

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<210> 1811
<211> 317
<212> nucleic acid
<213> Glycine max

<400> 1811

tgcacgcgta cgtaagctcg gaattcggct cgaggacacc tatggaggat ggggtgcaca 60
tgggtggtggt gccttctctg ggaaggatcc tatcaagggt gataggagtgt gtgcctacat 120
tgtgaggcaa gctgcaaaga gcattgttnc caaatggact tgctaggagg gcaattgtgc 180
aagtttctcta tgccattggt gtgcctgagc cttgtctgt gtttgttgac acttatggca 240
ctgggaagat ccctgacaag gaaatcctca gcattgtgaa ggagagtttt gactcaggcc 300
tggatgatct cnatcac 317

<210> 1812
<211> 323
<212> nucleic acid
<213> Glycine max

<400> 1812

acgtcncang cangcgtacg taagctcggn attcggctcg aggcttgaga aatggcacaa 60
ganacctttc tattcacatc tgaatctgta aacgagggtc accccgacaa gctgtgcgaa 120
ccagatctct gatgcagtgc tcgatgcgtg cttgaacag gaccctgaca gcaaggttgc 180
ctgtgagaca tgcaccaaga ccaacatggt catggtcttt ggagngatca canccaaggg 240
ccnnacgtag nctatgagaa gattgtccgt gacacctgcc gcgaaattgg attcatctct 300
gatgtgttcg gtcnngatgc gcc 323

<210> 1813
<211> 342
<212> nucleic acid
<213> Glycine max

<400> 1813

gctgcaaaga gtgttgtggc aaatggcctt gccagaaggt gcattgtcca agtttcctat 60
gccattggtg tccctgagcc cttgtcagtg tttgtggaca cttatggaac tgggaagatt 120

cctgacaagg agattottca aattgtgaag gagaatttcg acttcagacc tggaatgatc 180
 accattaact tggacottaa gaggggtggc cataggttcc tcaagacagc tgcttatgga 240
 cactttggaa gggatgacct gacttcacct gggaagttgt gaagccatca agtctgagaa 300
 gccncaactt agatgtgtga gttaaccatc ccttcattggn gc 342

<210> 1814
 <211> 318
 <212> nucleic acid
 <213> Glycine max
 <400> 1814

agtgcgancg acgcgtacgt aagctcgaa ttcggctcga gacggctgcg agaagcgaca 60
 gaaggggaag aaagatcatc attgatacct atggtggctg ggggtgctcat ggtggagggtg 120
 ccttttcagg gaaggacct accaaggttg acagaagtgg tgcctatatt gtaaggcagg 180
 ctgcaaagag tgtcgtggca aatggccttg ctagaagggtg catttgtgtca gtttcctatg 240
 ccattgggtgt ccctgagccc ttgtcagtgt ttgtggacac ttatggaaact gggaagattc 300
 ctgacaagga gattctgc 318

<210> 1815
 <211> 280
 <212> nucleic acid
 <213> Glycine max
 <400> 1815

cgcntgcacg cgtacgtaag ctcggaattc ggctcgagca cctatggagg atgggggtgca 60
 catggtgggtg gtgccttctc tgggaaggat cctaccaagg ttgataggag tgggtgcctac 120
 attgtgaggc aagctgcaaa gagcattgtt gcaaattggac ttgctaggag ggcaattgtg 180
 caagtttctt atgccattgg tgtgcctgag cccttgtctg tgtttgttga cacttatggc 240
 actgggaaga tccctgacaa ggaaatcctc agcattgtgn 280

<210> 1816
 <211> 236
 <212> nucleic acid
 <213> Glycine max
 <400> 1816

gagcattgtg gcaagtggac ttgccagaag gtgcattgtg caagtgtctt atgccattgg 60
 tgtgcctgag cctttgtctg tgtttgttga cacctatggc actgggaaga tccatgataa 120
 ggagattctc aacattgtga aggaaaactt tgatttcagg cctgggtatga tctccatcaa 180
 ccttgatctc aagaggggtg gaaataacag gtttttgaag actgctgcct atggac 236

<210> 1817
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 1817

acgtgcang cacgngacg taagctcgga attcggtcg aggtgccttc tctgggangg 60
 atcctaccaa ggttgatatg antggtgcct anattgtgag gcaagctgca aagagcattg 120
 ttgcaaattg acttgctagg agggcaattg tncaagtttc ctatgccatt ggtgngcctg 180
 agcccttntc tgtgtttgtt gacacggatg gcaactgggaa gatccctgac aangnaatcc 240
 tcagcattgt gaaggagagt ttgacttca ggctggcct gatctccatc naccttgagc 300
 tcaagagggg tggn 314

<210> 1818
 <211> 267
 <212> nucleic acid
 <213> Glycine max
 <400> 1818

gtgcganga cgcgtagctn agctcggaat tcggctcgag caatgacgaa attgctgctg 60
 acctcaaaga gcatgtgatc aagcctgtga tcccagagaa gtaccttgat gagangacca 120
 ttttccactt gaacccttca ggccgttttg tcattggtgg ccctcatggc gatgctggtc 180
 tcaccggccg caagatcatt atcgatactt atggaggatg gggtgctcat ggtggtggtg 240
 ctttctccgg gaaggaccct accaagg 267

<210> 1819
 <211> 278
 <212> nucleic acid
 <213> Glycine max
 <400> 1819

gtcgcangca cgcntacgta agctcggaaat tcggctogag ctctgggaag gatcctacca 60
 aggttgatag gagtgggtgcc tacattgtga ggcaagctgc aaagagcatt gttgcaaattg 120
 gacttgctag gagggcaatt gtgcaagttt cctatgccat tgggtgtgcct gagcccttgt 180
 ctgtgtttgt tgacacttat ggcaactggga agatccctga caaggaaatc ctcagcattg 240
 tgaaggagag ttttgacttc aggcctggca tgatctcc 278

<210> 1820
 <211> 281
 <212> nucleic acid
 <213> Glycine max

<400> 1820

catcgtatgc ncgcgtacgt aagctcggaa ttcggctcga gaattgtgca agtttcctat 60
 gccattggtg tgcctgagcc cttgtctgtg tttgttgaca cttatggcac tgggaagatc 120
 cctgcacaag gaaatcctca gcattgtgaa ggagagtttt gacttcaggc ctggcatgat 180
 ctccatcaac cttgatctca agaggggtgg aaatggcagg ttcttgaaga ctgctgcata 240
 tggacacttt ggcagagatg accctgactt cacatgggaa g 281

<210> 1821
 <211> 255
 <212> nucleic acid
 <213> Glycine max

<400> 1821

cttgccagaa ggtgcattgt gcaagtgtct tatgccattg gtgtncctga gcctttgtac 60
 tgtgtttgtt gacacctatg gcactggaaa gatccctgac aaggagntcc ttaacattgt 120
 gaaggagaac tttgatttca ggccctggtat gatctccatc aaccttgatc tcaagagggg 180
 nggaaataac aggtttttga agactgctgc atatggacac tttggaagag aggaccctgg 240
 acttcacatg ggaag 255

<210> 1822
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 1822

gtgcgcatgca cgcgtacgta agctcggaat tcggctcgan ggagtgggtgc ctacattgtg 60
 angcaagctg caaagagcat tgttgcaa at ggacttgcta ggagggcaat tgtgcaagtt 120
 tcctatgcc a ttggtgtgcc tgagcccttg tctgtgtttg ttgacctta tggcactggg 180
 aagatccctg acaaggaa at cctcagcatt gtgaaggaga gttttgactt caggctggca 240
 tgatctccat caacttgatc tcaagagggg tgggaatggc aggttcttga gatgctgcaa 300

<210> 1823
 <211> 283
 <212> nucleic acid
 <213> Glycine max

<400> 1823

gtgcgacgca cgcgtacgta agctcggaat tcggctcgag gtgcctacat tgtgaggcaa 60
 gctgcaaaga gcattgttgc aaatggactt gctaggaggg caattgtgca agtttcctat 120
 gccattggnc tgccctgagcc cttgtctgtg tttgttgaca cttatggcac tgggaagatc 180
 cctgacaagg aaatccctcag cattgtgaag gagagttttg acttcaggcc tggcatgatc 240
 tcatcaacct tgatctcaag aggggtggaa atggcaggtt ctt 283

<210> 1824
 <211> 306
 <212> nucleic acid
 <213> Glycine max

<400> 1824

accgtgnann cacnctacg taagctcgga attcggctcg agnggcaggt ttgtcattgg 60
 agggccgcat ggcgatgntg gtctcaccgg ccgcaagatc atcatcgaca cctatggagg 120
 atggggtgca catggtggtg gtgccttctc tgggaaggat cctaccaagg ntgataggag 180
 tgggtgcctac attgtgaggc aagctgcaaa gagcattggt gcaa atggac ttgctaggag 240
 ggcaattgtg caagtttctt atggccattg gtgtgcctga gcccttgtct gtgtttgtng 300
 acactt 306

<210> 1825
 <211> 313
 <212> nucleic acid
 <213> Glycine max

<400> 1825

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ggagggtggt gctggtgacc 60
agggtcacat gtttggctat gccactgatg aaaccctga gtacatgcc ctcagccatg 120
tccttgcaac caaactcggg gctcgctca ccgagggttag gaaaaatggt acctgtgctt 180
ggctgaggcc agatggcaag acacaagtaa ctgttgagta ctacaatgac aatggtgcca 240
tggttcagat tcgtgtccac antgtcntaa tttccacca ncatgatcct nctgtgagca 300
tgaccaaatt ggt 313

<210> 1826

<211> 357

<212> nucleic acid

<213> Glycine max

<400> 1826

ccaccacacc tctcctcggt cacgctaccc ctttctgctc ttcttctacc tttcaagttt 60
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 120
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 180
cccagacagc aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 240
gaaatcacga ccaaggccaa cggtgactac gagaagatag tgctgacac ctgcaggaac 300
atcggttcg tctcaaatga tgtgggatgg atgccgacaa ctgcaaggtc ctgctca 357

<210> 1827

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 1827

tgncgcatgc actcgtacgt aagctcgga ttcggctcga ggcaagggtc tcgtcaacat 60
tgagcaacag agtctgata ttgctcaagg tgtgcacggc caccttcaca aagaggcctg 120
aggagattgg tgctgggtgac caaggctcata tggtcggcta tgccactgac gagactcccg 180
agctcatgcc cttgagccat gtccttgcca cgaagctcgg tgccaagctc accgaggttc 240
ggaagaacgg gacatgccct tggtgagac ctgatggcaa gaccaagtc atgttgagta 300
tacaatgaca agggtgccat 320

<210> 1828
 <211> 282
 <212> nucleic acid
 <213> Glycine max

 <400> 1828

 nngtcgcatg cacgcgtacg taagctcgga attcggtctg agggctctcg tcaacattga 60
 gcaacagagt cctgatattg ctcaagggtg gcaaggccac ctcaaaaga ggcttgagga 120
 gattggtgct ggtgaccaag gtcatatgtt cggtatgcc actgacgaga ctcccgagct 180
 catgcccttg agccatgtcc ttgccacgaa gctcgggtgcc aagctcaccg aggttcggaa 240
 gaacgggaca tgcccttggc tgagacctga tggcaagacc ca 282

<210> 1829
 <211> 283
 <212> nucleic acid
 <213> Glycine max

 <400> 1829

 cgtgacacat gccgcgaaat tggattcatc totgatgatg ttggtottga tgctgacaaa 60
 tgcaagggtg tggtaacat tgagcagcag agccctgata tcgccagggt tgtgcacggg 120
 cacttcacca agcgcacaga ggagggtggt gctgggtgacc agggtcacat gtttggctat 180
 gccactgatg aaaccctga gtacatgcc ctcagccatg tottgcaacc aaactcgggtg 240
 ctgctcacc gaggttagga aaaatggtac tgtgcttggc tga 283

<210> 1830
 <211> 290
 <212> nucleic acid
 <213> Glycine max

 <400> 1830

 nncangcac gcgtacgtaa gctcggatt cggtcagagn cgtgacacct gcaggaacat 60
 cggcttcgtc tcaaatgatg tgggactgga tgccgacaac tgcaagggtc tcgtcaacat 120
 tgagcagcag agccctgata ttgctcagggtgtgtacacggc caccttacca aaaaacctga 180
 agaaattggt gctgggtgacc agggtcacat gtttggctat gccactgatg aaaccctga 240
 attgatgcc ttgagccatg ttcttgcaac aaaactcgggt gctcgtctca 290

<210> 1831
 <211> 268
 <212> nucleic acid
 <213> Glycine max

 <400> 1831

 gtcgcangca cgcgtacgta agctcggaat tcggctcgag catgcttgnc agcaggaccc 60
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atggatgatgg ttttcggaga 120
 gatcacaacc aaggccaacg tggactatga gaagattgtg cgtgacacat gcaggaacat 180
 tggttttgtc tctgatgatg ttggtcttga tgcgtgacaac tgcaaggtcc tcgtcaacat 240
 tgagcaacag agtcctgata ttgctcaa 268

<210> 1832
 <211> 315
 <212> nucleic acid
 <213> Glycine max

 <400> 1832

 cgcangcacg cgtacgtaag ctcggaattc ngctcgangg agattgggtgc tggtgaccaa 60
 ggtcatgggt tcggctatgc cactgacgag actcccgagc tcatgccctt gagccatgtc 120
 cttgccacga agctcgggtgc caagctcacc gaggttcgga agaacgggac atgcccttgg 180
 ctgagacctg atggcaagac ccaagtcact gttgagtact acaatgacaa gggtgccatg 240
 gttccaatcc gcgtccacac tgtgtcatc tncacacaac atacgaccng agtgtggncg 300
 cggattggna catgg 315

<210> 1833
 <211> 240
 <212> nucleic acid
 <213> Glycine max

 <400> 1833

 agaaattggg gctggtgacc agggtcacat gtttggctat gccactgatg aaaccctga 60
 attgatgcca ttgagccatg ttcttgcaac aaaactcggg gtcggtctca ccgaggttcg 120
 caagaacggg acctgccctt ggctgaggcc tgatgggaag acccaagtga ccgttgagta 180
 ttacaatgac aatggtgcca gggttcctat tcgtgtacac accgtgctaa tctccacca 240

<210> 1834
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 1834

 antcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggctcgatgc atgcttggag 60
 caggaccctg acagcaaggt tgctgtgaa acctgncacc aagaccaaca tggatgatggt 120
 tttcggagag atcacaacca aggccaacgt ggactatgag aagattgtgc gtgacacatg 180
 caggaacatt ggtccaagtc totgatgatg ttggtcttga tgcgtgacaac tgcaagggtcc 240
 tcgtcaacat tgagcaacag agtcctgata ttgctcaagg tgtgcacggc cacctc 296

<210> 1835
 <211> 286
 <212> nucleic acid
 <213> Glycine max

 <400> 1835

 gtcgcatgca cgcgtacgta agctcggaa ttcggctcga gcaagatcat tatcgatact 60
 tatggaggat ggggtgctca tgggtggtgt gctttctccg ggaaggaccc taccaagggt 120
 gataggagtg gtgcttacat tgtgagacag gctgctaaga gcatgtggca agtggacttg 180
 ccagaagggt cattgtgcaa gtgtcttatg ccattggtgt gcttgagcct ttgtctgtgt 240
 ttgttgacac ctatggcact gggaagatcc atgataagga gattct 286

<210> 1836
 <211> 341
 <212> nucleic acid
 <213> Glycine max

 <400> 1836

 nacangcacg cntacgtaag ctccggaattc ggctcgagct caagtttttg aagtatagag 60
 atggcagaga cattcctatt tacctcagag tcagtgaacg agggacaccc tgnccaagct 120
 ctgtgaccaa atctctgatg ctgtcctcga cgcttgccctc gaacaggacc cagacagcaa 180
 ggttgccctgc gaaacatgca ccaaaaccaa cttgggtcatg gtcttcggag aaatcacgac 240
 caaggccaat gttgactacg agaagatagt gcgtgacacc tgcaggaaca tcggctttgt 300

ctcaaacgat gtgggactgg atgccgacaa tgcaaggtcc t 341

<210> 1837
 <211> 313
 <212> nucleic acid
 <213> Glycine max

<400> 1837

gtcnnangca cgntacgtaa gctcggaatt cggctcganc tcgagccgaa tcggctcgag 60
 gccgcatggc gatgctggtc tcnncggcc gcaagatcan catcgacacc tatggaggan 120
 ggggtgcaca tgggtggtgt gccttctctg ggaaggatcc taccaagggt gataggagtg 180
 gtgcctacat tgtgaggcaa gctgcaaaga gcattgttgc aaatggactt gctaggaggg 240
 caattgtgca agtttcttat gccattggtg tgctgagcc cttgtctgtg tttgttgaca 300
 cttatggcac tgg 313

<210> 1838
 <211> 276
 <212> nucleic acid
 <213> Glycine max

<400> 1838

nangcacgcg tacgtaagct cggaattcgg ctcgaggaca cctgcaggaa catcggcttc 60
 gtctcaaattg atgtgggact ggatgccgac aactgcaagg tcctcgtcaa cattgagcag 120
 cagagccctg atattgctca ggggtgtacac ggccacctta ccaaaaaacc tgaagaaatt 180
 ggtgctggtg accagggtca catgtttggc tatgccactg atgaaacccc tgaattgatg 240
 ccattgagcc atgttcttgc aacaaaactc ggtgct 276

<210> 1839
 <211> 286
 <212> nucleic acid
 <213> Glycine max

<400> 1839

angtcgcang cacgcgtagc taagctcgga attcggctcg agntgcacca aaaccaactt 60
 ggtcatggtc ttcggagaaa tcacgaccaa ggccaacgtt gactacgaga agatagtgcg 120
 tgacacctgc aggaacatcg gcttcgtctc aaatgatgtg ggactggatg ccgacaactg 180

caaggtcctc gtcaacattg agcagcagag ccttgatatt gctcagggtg tacacggcca 240
ccttaccaaa aaacctgaag aaattggtgc tggtgaccag ggtcac 286

<210> 1840
<211> 315
<212> nucleic acid
<213> Glycine max

<400> 1840

gtcgctagca cgcgtacgta agctcggaat tcggctcgag ntttctatg ccattggtgt 60
ccctgagccc ttgncagtgt ttgtggacac ttatggaact gggaagattc ctgacaagga 120
gattcttcaa attgtgaagg agaatttoga cttcagaact ggaatgatca ccattaactt 180
ggaccttaag aggggtggcc ataggttctt caagacagct gcttatggac actttggaag 240
ggatgaccct gacttcacct gggaagttgt gaagccactc aagtctgaga agcctcaagc 300
ntaagattgt tgtga 315

<210> 1841
<211> 408
<212> nucleic acid
<213> Glycine max

<400> 1841

gagaagacga cagaaggggg cagcgcttga tttgaggcca ggcaagcccc actcaaccac 60
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agtataaaga tggcagagac attcctatth acctcagagt cggatgaacga gggacaccct 180
gacaagctct gcgaccaaht ctccgatgct gtctctgacg cttgcctcga gcaggaccca 240
gacagcaaag ttgcctgcga aacatgcacc aaaaccaact tggatcatggt cttcggagaa 300
atcacgacca aggccaacgt tgactacgag aagatagtgc gttacacctg caagaacatc 360
cgntttctct cnaatgattt ggaactggat nccaaaaatt gcaaggte 408

<210> 1842
<211> 255
<212> nucleic acid
<213> Glycine max

<400> 1842

ttgataccta tgggtggctgg ggtgctcatg gtggaggtgc cttttcaggg aaggacccta 60
ccaaggttga cagaagtggg gcctatatattg taaggcaggc tgcaaagagt gtcgtggcaa 120
atggccttgc tagaaggtgc attgtgcaag tttcctatgc cattgggtgtc cctgagccct 180
tgtcagtgtt tgtggacact tatggaactg ggaagattcc tgacaaggag attctgcaat 240
tgtgaaggag attcc 255

<210> 1843
<211> 273
<212> nucleic acid
<213> Glycine max
<400> 1843

tctcaagttt ttgaagtata gagatggcag agacattcct atttacctca gagtcagtga 60
acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 120
tcgaacagga ccagacagc aaggttgcct gcgaaacatg caccaaaacc aacttggtca 180
tggtcttcgg agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca 240
cctgcaggaa catcggttt gtctcaaacg atg 273

<210> 1844
<211> 272
<212> nucleic acid
<213> Glycine max
<400> 1844

nacgcangcn cgcgtacgta agctcggaat tcggctcgag agtggacttg ccagaagggtg 60
cattgtgcaa gtgtcttatg ccattgggtg gcctgagcct ttgtctgtgt ttgttgacac 120
ctatggcact gggaagatcc atgataagga gattctcaac attgtgaagg aaaactttga 180
tttcaggcct ggtatgatct ccatcaacct tgatctcaag aggggtggaa ataacaggtt 240
tttgaagact gctgcctatg gacactttgg aa 272

<210> 1845
<211> 279
<212> nucleic acid
<213> Glycine max
<400> 1845


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gtcgcangca cgcgtacgta agctcggaat tcggctcgag cacgaagctc ggtgccaagc 60
tcaccgaggt tcggaagaac gggacatgcc cttggctgag acctgatggc aagacccaag 120
tcactgttga gtactacaat gacaaggggtg ccatggtncc caatccgctg ccacactgtg 180
ctcatctcca cacagcatga tgagactgtc acaaatgatg agattgcagc tgatcttaaa 240
gaacacgtga ttaagcctgt gattcctgag aagtacctt 279

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<210>      1846
<211>      269
<212>      nucleic acid
<213>      Glycine max

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<400>      1846

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gtcgcattgca cgcgtacgta agctcggaat tcggctcgag gaccaacatg gtgatggttt 60
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<210>      1847
<211>      439
<212>      nucleic acid
<213>      Glycine max

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<400>      1847

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ggcaagcccc actcaaccac cacacctctc ctcggtcacg ctaccccttt ctgctcttct 120
tctacctttc aagtttttaa agtataaaga tggcagagac attcctattt acctcagagt 180
cgggtgaacga gggacaccct gacgagctct gcgaccaa atccgatgct gtcctcgacg 240
cttgccctga gcaggacca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 300
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gcaaggtcct cgtcaacat 439

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<210>      1848

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<211> 407
 <212> nucleic acid
 <213> Glycine max

<400> 1848

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tcgaggccag gcaagcccca ctcaaccacc acacctctcc tcgttcacgc tacccttttc 60
tgctcttctt ctacctttca agttttaaaa gtataaagat ggcagagaca ttctatttta 120
cctcagagtc ggtgaacgag ggacaccctg acaagctctg cgaccaaata tccgatgctg 180
tctcgcacgc ttgcctcgag caggaccctg acagcaaagt tgcttcgcga acatgcacca 240
aaaccaactt ggtcatggtc ttccggagaaa tcacgaccaa ggccaacgtt gactacgaga 300
agatagtgcg tgacacctgc aagaacatcg ggttcgtccc aaatgatgtt tggaactggg 360
gttccgacaa ctgggaaagg ttctcgttca anattgagca agcaaag 407
```

<210> 1849
 <211> 282
 <212> nucleic acid
 <213> Glycine max

<400> 1849

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cagtgaacga gggacaccct gacaagctct gtgaccaaata ctctgatgct gtcctcgacg 120
cttgctcga acaggaccca gacagcaagg ttgcctgcga aacatgcacc aaaaccaact 180
tggtcatggt ctccggagaa atcacgacca aggccaatgt tgactacgag aagatagtgc 240
gtgacacctg caggaacata ggctttgtct caaacgatgt gg 282
```

<210> 1850
 <211> 266
 <212> nucleic acid
 <213> Glycine max

<400> 1850

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cattcctgag aagtacctgg atgagaagac catcttccac cttaaccctt ctggccgttt 120
tgtcattggt ggccctcatg gtgatgctgg tctcactgga agaaagatca tcattgatac 180
ctatggtggg tggggtgctc atggtggagg tgccttttca gggaaggacc ctaccaaggt 240
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tgacagaagt ggtgcctata tcgtga

266

<210> 1851
<211> 272
<212> nucleic acid
<213> Glycine max

<400> 1851

ggtgctcatg gtggtggtgc cttctccggg aaggatccca ccaaggttga taggagtgg 60

gcttacattg tgagacaggc tgctaagagc attgtggcaa gtggactagc cagaaggtgc 120

attgtgcaag tgtcttatgc cattgggtgtg cccgagcctt tgtctgtctt tgttgacacc 180

tatggcaccg ggaagatcca tgataaggag attctcaaca ttgtgaagga gaatttgatt 240

ncaggcccgg tatgatctcc atcaaccttg at 272

<210> 1852
<211> 305
<212> nucleic acid
<213> Glycine max

<400> 1852

cnncatncgt aagtnantnc nncattnggc tcgagccaac ttggtcatgg tctncggata 60

natcntgncc aaggccancg ttgnctacga gnagatagtg cgtgacacct gcaggaacat 120

cggcttcgtc tcanatgatg tgggactgga tgccgacaac tgcaaggtcc tcgtcaacat 180

tgagcagcag agccctgata ttgctcaggg tgtacacggc caccttacca aaaaacctga 240

agaaattggg gctggtgacc agggtcacat gtttggctat gccatgatga nccctgaatt 300

gatgc 305

<210> 1853
<211> 340
<212> nucleic acid
<213> Glycine max

<400> 1853

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tgttcaagct accatctctc tctctctttc ttagtgctc cttgccagaa gttaaaatgg 120

cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180

gtgaccagat ctccgatgct gtgctcgatg catgcttga gcaggaccct gacagcaagg 240
 ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca 300
 aggccaacgt ggactatgag aagattgtgc gtgacacatg 340

<210> 1854
 <211> 329
 <212> nucleic acid
 <213> Glycine max
 <400> 1854

agtcgcangc acgcntacgt aagctcggaa ttcggctoga gctctctgtt ctcttctacc 60
 tctcaagtnt ttgaagtata gagatggcag agacattcct atttacctca gactcngtga 120
 acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 180
 tcgaacagga cccagacagc aaggttgccct gcgaaacatg caccaaaacc aacttggtca 240
 tggctctcgg agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca 300
 cctgcaggaa catcggttt gtctcaaac 329

<210> 1855
 <211> 293
 <212> nucleic acid
 <213> Glycine max
 <400> 1855

tcgcatgcac gcntacgtaa gctcggatt cggtcgcagn ganattggtg ctggtgacca 60
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 tcttgcaaca aaactcgggtg ctctgtcac cgaggttcgc aagaacggtg cctgcccttg 180
 gctgaggcct gatgggaaga cccaagtgc cgttgagtat tacaatgaca atggtgccag 240
 ggttcctatt cgtgtacaca ccgtgcnaa tctccacca acacgacgag nct 293

<210> 1856
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 1856

naaangtcgc atgcacgcgt acgtaagctc ggaattcggc tcgagtgcac gttgagtatt 60

acaatgacaa tgggtgccagg gttcctattc gtgtacacac cgtgctaata tccacccaac 120
acgacgagtn ctgtcaccaa tgacgaaatt gctgctgacc tcaaagagca tgtgatcaag 180
cctgtgatcc cagagaagta ccttgatgag aagaccattt tccaacttga acccttcagg 240
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cgatac 306

<210> 1857
<211> 294
<212> nucleic acid
<213> Glycine max
<400> 1857

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ggtcacatgt ttggctatgc cactgatgaa acccctgagt acatgccct cagccatgtc 120
cttgcaacca aactcgggtgc tcgcctcacc gaggttagga aaaatggtag tgtgcttggc 180
tgaggccaga tggcaagaca caagtaactg ttgagtacta caatgacaat ggtgccatgg 240
ttccagttcg tgtccacact gtcttaattt ccacacaaca tgnnnnnacc ganc 294

<210> 1858
<211> 394
<212> nucleic acid
<213> Glycine max
<400> 1858

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cacacctctc ctggttcacg ctaccccttt ctgctcttct tctaccttct aagttttaaa 120
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agacagcaaa gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg tcttcggaga 300
aatcacgacc aaggccaacg ttgactacga gaagatagtg cgtgacacct gcangaacat 360
cggcttcgtc tcaaatgatg tgggactgga tgcc 394

<210> 1859
<211> 307
<212> nucleic acid

<213> Glycine max

<400> 1859

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tgctgacaaa tgcaagggtg ttgtcaacnt tgagcagcag agccctgata tcgcccaggg 120
tgtgcacggt cactttcacc aagcgcccag aggagggttg tgctgggtgnc cagggtcaca 180
tgtttggtta tgccactgat gaaacccctg agtacatgcc cctcagccat gtccttgcaa 240
ccaaactcgg tgctcgctc accgagttag gaaaaatggt acctgtgctt ggctgaggcc 300
agatngc 307

<210> 1860

<211> 493

<212> nucleic acid

<213> Glycine max

<400> 1860

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gctgcgaaga aaacgacaga aggggggcagc gcttgatttg aggccaggca agccccactc 120
aaccaccaca cctctcctcg ttcacgctac ccctttctgc tcttcttcta cctttcaagt 180
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 240
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 300
gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggc catggtcttc 360
ggagaaatca cgaccaangg caacgttgac tacnanaann aaattncntg acacctgcag 420
gaacatcggc ttctgtctca atgatgtggg actgggatgc cgacaactgc aangtcctcg 480
tcaacattga gca 493

<210> 1861

<211> 489

<212> nucleic acid

<213> Glycine max

<400> 1861

ggcttttnng ccngtnnaa tcttacagn caggtaccgg tacggaattc ccggctcgac 60
ccacgcgtac gtacggctgc gagaagacga cagaaggggg cagcgcttga tttagggcca 120

ggcaagcccc actcaaccac cacacctctc ctcgttcacg ctaccccttt ctgctcttct 180
 tctacctttc aagttttaaa agtataaaga tggcagagac attcctattt acctcagagt 240
 cggtgaacga gggacaccct gacaagctct gcgaccaa atctcgatgct gtctctgacg 300
 cttgcctcga gcaggacca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 360
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 ccttaacacc tgcaggaaca tcggttctgt ctcaa atgat gtgggactgg atgccgacaa 480
 ctgcaaggt 489

<210> 1862
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 1862
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 gccctgatat tgctcagggg gtacacggcc accttacc aaacacgtgaa gaaattgggt 180
 ctggtgacca gggtcacatg ttgggtatg ccaatgatga aaccctgaa ttgatgccat 240
 tgagccatgt tcttgcaaca aaactcgggt ctcgtctcac cgagggttcgc aagaacggta 300

<210> 1863
 <211> 330
 <212> nucleic acid
 <213> Glycine max

<400> 1863
 ncatgcacgc gtacgtaagc tcggaattcg gctcgagctt ctacctctca agtttttgaa 60
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 gacaagctct gtgaccaa atctctgatgct gtctctgacg cttgcctcga acaggacca 180
 gacagcaagg ttgcctgcga aacntgcacc aaaaccaact tggatcatggt cttcggagaa 240
 atcacgacca aggccaatgt tgactacgag aagatagtagc gtgacacctg caggaacatc 300
 ggctttgtct caaacgatgt gggactggat 330

<210> 1864

<211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 1864

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 ttccggctcga gcatgggtgga gntgcctttt cagggaaagn ccctaccaag gttgacagaa 120
 gtgggtgccta tatcgtgagg caggtcgcaa agagtgttgt ggcaaattggc cttgccagaa 180
 ggtgcattgt ccaagtttcc tatgccattg gtgtccctga gcccttgtca gtgtttgtgg 240
 acacttatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 300
 cgacttca 308

<210> 1865
 <211> 288
 <212> nucleic acid
 <213> Glycine max

 <400> 1865

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 aagtttccta tgccattggt gtccctgagc ccttgtcagt gtttgtggac acttatggaa 120
 ctgggaagat tcctgacaag gagattcttc aaattgtgaa ggagaatttc gacttcagac 180
 ctggaatgat caccattaac ttggacctta agaggggtgg ccataggttc ctcaagacag 240
 ctgcttatgg acactttgga agggatgacc ctgacttcac ctgggaag 288

<210> 1866
 <211> 281
 <212> nucleic acid
 <213> Glycine max

 <400> 1866

 gcgtacgtna gctcggaatt cggctcgagn gcaagtttcc tatgccattg gtgtccctga 60
 gcccttgtca gtgtttgtgg acacttatgg aactgggaag attcctgaca aggagattct 120
 gcaaattgtg aaggagaatt tcgacttcag acctggaatg atcaccatta acttggacct 180
 taagaggggt ggtcataggt tcctcaagac agctgcttat ggacactttg gaagggatga 240
 tgcagacttc acctgggaag ttgtgaagcc actcaagtca g 281

naangnagan gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gtttttgaag 60
tatagagatg gcagagacat tcctatttcc ctcagagtca gtgaacgagg gacaccctga 120
caagctctgt gaccaaactct ctgatgctgt cctcgacgct tgcctcgaac aggaccaga 180
cagcaagggtt gcctgcgaaa catgcaccaa aaccaacttg gtcatgggtct tcggagaaat 240
cacgaccaag gccatgttga ctacgagaag atagtgcgtg acacctgccg gaacatcggc 300
tttgtctcna acgatgtggg at 322

<210> 1870
<211> 418
<212> nucleic acid
<213> Glycine max

<400> 1870

tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 60
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 120
aagctctgtg accagatctn cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 180
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acaaccaagg ccaacgtgga ctatgagaaa gattgtgcgt gacacatgca ggaaccattg 300
ggttttgctc tgatgaatgt ggtcttggat gcttgacact gcaaggctcct cgtcaacatt 360
tgagcaacag aagtctgat antgnttcaa ggtgtgcacg ggcacottac aaaagang 418

<210> 1871
<211> 261
<212> nucleic acid
<213> Glycine max

<400> 1871

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tgcctcgaac aggaccaga cagcaagggtt gcctgcgaaa catgcaccaa aaccaacttg 180
gtcatgggtct tcggagaaat cacgaccaag gccaatgttg actacgagaa gatagtgcgt 240
gacacctgca ggnacatcgg g 261

<210> 1872

<211> 277
 <212> nucleic acid
 <213> Glycine max

<400> 1872

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 gcaagggttg ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 180
 caaccaaggc caacgtggac tatgagaaga ttgtgcgtga cacatgcagg aacattgggt 240
 ttgtctctga tgatgttggt cttgatgctg acnactg 277

<210> 1873
 <211> 291
 <212> nucleic acid
 <213> Glycine max

<400> 1873

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 actgcaaggt cctcgtcaac attgagcann cagagccctg atattgctca ggggtgtacac 180
 ggccacctta ccaaaaaacc tgaagaaatt ggtgctggtg accaggggtca catgtttggc 240
 tatgccactg atgaaacccc tgattgatgc cattgagcca tgttcttgca a 291

<210> 1874
 <211> 287
 <212> nucleic acid
 <213> Glycine max

<400> 1874

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 gtgatggttt tcggagagat cacaaccaag gccaacgtgg atatgagaag atgtgcgtga 240
 cacatgcagg aacattgggt ttgctctgat gaggttggctt gatgctg 287

<210> 1875

<211> 262
 <212> nucleic acid
 <213> Glycine max

<400> 1875

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 ttgtgcaagt ttctatgcc attggtgtgc ctgagccctg tctgtgtttg ttgacactta 180
 tggcactggg aagatccctg acaaggaaat cctcagcatt gtgaaggaga gttttgactt 240
 caggcctggc atgatctcca tc 262

<210> 1876
 <211> 315
 <212> nucleic acid
 <213> Glycine max

<400> 1876

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 tgcctgtgan acctgnanca agaccaacat ggtgatggtt tncngagaga tcanaaccaa 180
 ggccaacgtg gactatgaga agattgtgcg tgacacatgc aggaacattg gttttgtctc 240
 tgatgatggt ggtctgatgc tgacaatgca nagtcctcgt caacattgag cnacagagtc 300
 ctgatattgc tcaag 315

<210> 1877
 <211> 489
 <212> nucleic acid
 <213> Glycine max

<400> 1877

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cggcttccgt ctcaaacgat gtgggacttg atgctgacaa ctgcaanggt cttgtnaaca 480
ttgancacc 489

<210> 1878
<211> 468
<212> nucleic acid
<213> Glycine max
<400> 1878

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<210> 1879
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 1879

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atggtgccat ggttccagtt cgtgtccaca ctgtcctaatt ttccacccaa catgatgnga 240
cgtgagcaat gatcaaattg ctgcggacct taaagagcat gttatcaagc ctgtcattcc 300

<210> 1880
<211> 477
<212> nucleic acid

661

<213> Glycine max

<400> 1880

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 ttcggagaaa tcacgaccaa ggccaacgtt gactacgaga agatagtgcg tgacacctgc 420
 aagaacaacc ggttttctccc naattgaatn tgggactgga tgccgacaac tgcaang 477

<210> 1881

<211> 259

<212> nucleic acid

<213> Glycine max

<400> 1881

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 gtcacaaatg ntgagattgc agctgatctt aaagaacacg tgattaagcc tgtgattcct 180
 gagaagtacc ttgatgagag accattttcca tttgaaccct tccnggcagg ttgtcattgg 240
 agggcgggcat ggggatgng 259

<210> 1882

<211> 254

<212> nucleic acid

<213> Glycine max

<400> 1882

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 ntggttgagg cctgatggga agaccaagt gactgttgag tattacnntg acaacgggtgc 180
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254

<400> 1883

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<400> 1884

[illegible]

<400> 1885

662

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<210> 1886
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 1886

nntaannntn agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gaaagtataa 60
 agatggcaga gacattccta tttaacctcag agtcgggtgaa cgagggacac cctgacaagc 120
 tctgcgacca natctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca 180
 aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaaatcacga 240
 ccaaggccaa cgttgactac gagaagatag tgcgtgacac ctgcaggaac atcggtctcg 300
 t 301

<210> 1887
 <211> 508
 <212> nucleic acid
 <213> Glycine max
 <400> 1887

tnnnnnaact nagttctccg cngcctgna angtaagan gncccgggcc gacccacgcg 60
 tccntacggc tgcgagaaga cgacagaagg gggcagcgct tgatttgagg ccaggcaagc 120
 cccactcaac caccacacct ctctcgttc acgctacccc tttctgctct tttctacct 180
 ttcaagtttt aaaagtataa agatggcaga gacattccta tttaacctcag agtcgggtgaa 240
 cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt 300
 cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat 360
 ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gagaagatag tgcgtgacac 420
 ctgcangaac atcggtctcg tctcaaaatg atgtnggact ggattccac aactgcaaag 480
 ncctccnta aaattttgcc anaanccc 508

<210> 1888
 <211> 278
 <212> nucleic acid

<213> Glycine max

<400> 1888

gncgcacgcg tacgtaagct cggaattcgg ctcgagggca ggtttgtcat tggagggccg 60
catggcgatg ctggtctcac cgcccgcaag atcatcatcg acacctatgg aggatggggg 120
gcacatgggtg gtgggtgncct tctctgggaa ggatcctacc aagggttgnta ggagtgggtgc 180
ctacattntg aggcaagctg caaagagcat tgttgcaaat ggacttgcta ggagggcaat 240
tgtgcaagtt tcctatgcc a ttgggtgtgcc tgagccct 278

<210> 1889

<211> 280

<212> nucleic acid

<213> Glycine max

<400> 1889

cnttggtga gacctgatgg caagacccaa gtcantgttg agtactacaa tgacaagggt 60
gccatggttc caatnccggt ccacantgtg cttcatntnc acacagcatg atgagtgngt 120
nanaaatgat gagattgcag ctgatcttaa agaacacgtg attaagcntg tgatttctga 180
gaagtacctt gatgagaaga ccattttcca tttgaacct tntgggcagg tttgtcatgg 240
agggccgcag ggcgattttg gtgtnanggc ngnaagatcc 280

<210> 1890

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1890

noncangcnc gcnnncgtga gctcgnatt cggctcgagg tttttgangt atagagatgg 60
cagagacatt cctatttacc tcagagtcag tgaacgagg acacctgan aagctctgtg 120
accaaattctc tgatgctgtc ctgcacgctt gcctcgnaca ggaccagac agcaagggtg 180
cntgcgaaac atgcacaaaa accaacttgg tgcattggtct tcggagaaat caccaccaag 240
gccaatgttg actacgagaa gatagtgcgt gacacctgca ggaacatcgg ctttgtctca 300
aacgatgtgg 310

<210> 1891

<211> 290
 <212> nucleic acid
 <213> Glycine max

<400> 1891

cagcangcac gcgtacgtaa gctcggaatt cggctcgaga ccaacttggn atggtcttcg 60
 gagaaatcac gaccaaggcc aacgttgact acgagaagat agtgcgtagac acctgcagga 120
 acantcggct tcgtctcaaa tgatgtggga ctggatgccg acaactgcaa ggtcctcgtc 180
 aacattgagc agccgagccc tgatattgct caggggtgtac acggccacct taccaaaaaa 240
 cctgaagaaa ttgggtgctgg tgaccagggt cacatgtttg gctatgccat 290

<210> 1892
 <211> 502
 <212> nucleic acid
 <213> Glycine max

<400> 1892

tttaaactct ccgcggtcag gtaacggctn gngaattccc gggncgaccc angcgtccan 60
 nccaggcgtc agcccaacgc gtongtgngg ctgcnagaag annacanaag ggggcagcgc 120
 ttgatttnag gccaggcang cccactcat ccancanacc tctcctcgtt cangtncct 180
 ctttctgccc ttontctacc tticangttt taaaagtata nagatggcag agacattcct 240
 atttacctca gagtcggtga acgagggaca ccctgacaag ctctgctacc aaatctccga 300
 tgctgtcctc gacgcttgcc tcgagcanga ccagacagc naagttgcct gcgaaacatn 360
 caccatancc aacttggtca tgggtcttcgg aganatcacg accaaggcca acgttgacta 420
 cgaagaagat agtgcgtagac acctgcagga acatcngntt cgtctcaaata tatgtgggac 480
 tggatgccan canctgcaag gt 502

<210> 1893
 <211> 286
 <212> nucleic acid
 <213> Glycine max

<400> 1893

tcgcntgcac gcgtacgtaa gctcggaatt cggctcgagn aagttgcctg cganacatgc 60
 accaanacca acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 120

gagaagatag tgcgtgacac ctgcaggaac atcggttcg tctcaaata tgtgggactg 180
 gatgccgaca actgcaaggt cctcgtcaac attgagcngc agagccctga tattgctcag 240
 ggtgtacacg gccaccttac caaaaaacct gaagaaannc ntgctg 286

<210> 1894
 <211> 326
 <212> nucleic acid
 <213> Glycine max
 <400> 1894

ncgcngcacg cgtacgtaag ctcggaatc ggctcgagca caaagcgggt tactgtctgt 60
 tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt aaaatggccc 120
 aagaaaacttt cctattcaca totgaatcag tgaacgaggg gcaccctgac aagctctgtg 180
 accagatctc cgatgctgtg ctgatgcat gcttgagca ggaccctgac agcaagggtg 240
 cctgtgaaac ctgcaccaag accaactggt tgatgggttt cggagagatc acaaccaagg 300
 ccaacgtgga ctatgagaag attgtg 326

<210> 1895
 <211> 304
 <212> nucleic acid
 <213> Glycine max
 <400> 1895

tcgnangcaa gcncngaant cngcncnagc gnnnntagnc nanngtgntt accggcngca 60
 agatcantat cgatacttat nggaggatgn ngtgctcatg gtngtggtgc tttctccggc 120
 aaggacccta ccaaggttga taggagtggg gcttacattg tgagacaggc tgctaaganc 180
 attgtggcaa gtggactngc cagaaggtgc attgtgcaag tgtcttatgc cattggtgtg 240
 cctgagcctt tgtctgtgtt tgttgacacc tatggcactg ganagatccc tgacaaggag 300
 atct 304

<210> 1896
 <211> 273
 <212> nucleic acid
 <213> Glycine max
 <400> 1896

ncgtcgcatg cacgcgtacg tnagctcgga attcggtcgc agctgtgctt ggctgaggcc 60
 agatggcaag acacaagtaa ctgttgagta ctacaatgac aatgggtgcca tggttccagt 120
 tcgtgtccac actgtcctaa tttccacca acatgatgac nangtgagca atgaccaa 180
 tgctgtgac cttaaagagc atgttatcaa gctgtcatt cctgagaagt acctggatga 240
 gaagaccatc ttccacctta acccttctgg ccg 273

<210> 1897
 <211> 334
 <212> nucleic acid
 <213> Glycine max
 <400> 1897

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag cagcacaaag cgggttactg 60
 ncctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120
 tggcccaaga aactttccta ttcacatctg aatcagtga cgaggggac cctgacaagc 180
 tctgtgacca gatctccgat gctgtgctgc atgcatgctt ggagcaggac cctgacagca 240
 aggttgctg tgaaacctgc accaagacca acatggtgat ggttttcgga gagatcacia 300
 ccaaggccaa cgtggactat gagaagattg tgcg 334

<210> 1898
 <211> 293
 <212> nucleic acid
 <213> Glycine max
 <400> 1898

gtcgngcac gcgtacgtaa gctcggaatt cggctcgagn cctcacagag gttaggaaga 60
 atggcacctg tgcttggttg aggccagatg gtaagacaca agtaaccgtc gactactaca 120
 atgacaatgg tgccatggtt ccagttcgtg tccacactgt cctaatttcc acccaacatg 180
 acgacctgtg agccatgatc aaattgctgc ggaccttaaa gancatgtta tcaagcctgt 240
 cattcctgag aagtaccttg atgagaagac catcttccac ttaacccttc tgg 293

<210> 1899
 <211> 316
 <212> nucleic acid
 <213> Glycine max

<400> 1899

cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 60
tataaagatg gcagagacat tectattttac ctcagagtcg gtgaacgagg gacaccctga 120
caagctctgc gaccaaattct cccgatgctgt cctcgacgct tgcctcgagc aggaccacaga 180
cancaaagtt gcctgcgaaa catgcaccaa aaccaacttg gttcatggtc ttcgagagaaa 240
tcacgaccaa ggccaacggt gactacgaga agatagtgcg tgacacctgc aggaacatcg 300
gcttcgtctc aaatga 316

<210> 1900

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 1900

ttcctattta cctcagagtc ggtgaacgag ggacaccctg acaagctctg cgaccaaata 60
tcgatgctg tctacgacg cttgcctcga gcaggacca gacagcaaag ttgcctgcga 120
aacatgcacc aaaaccaact tggatcatgg cttcgagaaa atcacgacca aggccaacgt 180
tgactacgag aagatagtgc gtnacactgg agganacatg ggcttcgtct naaatgntgn 240
gggactggat cccganaant gaaggtcncg aaaatntga 279

<210> 1901

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1901

cgctgcacgc gtacgtaagc tcggaattcg gctcgagatg ctgtcctcga cgcttgctc 60
gagcaggacc cagacagcaa agttgcctgc gaaacatgca ccaaaccaca cttggatcatg 120
gtcttcggag aaatcacgac caaggccaac gttgactacg agaagatagt gcgtgacacc 180
tgcaggaaca tcggcttcgt ctcaaatgat gtgggactgg atgccgacaa ctgcaagggt 240
cctcgttcaa cattgagcag cagagccctg atattgctca ggggtg 285

<210> 1902

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 1902

gtcgcacatgca cgcgtacgta agctcgggaat tcggctcgag caggacccag acagcaaagt 60
tgcttgcgaa acatgcaaaa accaacttgg tcatgggtctt cggagaaatc acgaccaagg 120
ccaacgttga ctacgagaag atagtgcgtg acacctgcag gaacatcggc ttcgtctcaa 180
atgatgtggg actggatgcc gacaactgca aggtcctcgt caacattgag cagcagagcc 240
ctgatattgc tcaggggtgta cacggccacc ttaccaaaaa ac 282

<210> 1903

<211> 476

<212> nucleic acid

<213> Glycine max

<400> 1903

tttactnnnc cngngccatg taanagtana gaagtcccgg gccgaccac gngtcnntac 60
ggctgcgaga agacgacaga agggggcagc gcttgatttg aggccaggca agccccactc 120
aaccaccaca cctctctcgt ttcacgttac cctttctgc tctttctcna cctttcaagt 180
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 240
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcat 300
gaccagaca gcaaagttgc ctgcgaaaca tgcaccagaa ccaacttggc catgggtcttc 360
ggagaaatca cgaccatggg caacgttgac taccagaaga taagtgcgtg acacctgcag 420
gaacatcggg ttcgtctcaa atgatgtggg actggatgcc nacaattcgg anangg 476

<210> 1904

<211> 496

<212> nucleic acid

<213> Glycine max

<400> 1904

aactttcgtg ccagnccggt caagaatncc gggtcgaccc acgcgtccgt acggctgcga 60
gaagacgaca gaagggtacg gctgcgagaa gacgacagag ggggcagcgc ttgatttgag 120
gccaggcaag cccactcaa ccaccacacc tctcctcgtt cacgtaccc ctttctgctc 180
ttcttctacc tttcaagttt taaaagtata aagatggcag agacattcct atttacctca 240

gagtcggtga acgagggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc 300
gacgcttgcc tcgagcagga cccagacagc aaagttgctt gcgaaacatg caccaaaaacc 360
aacttggtca tggctcttcgg agaaatcacg accaaggcca acgttgacta cgaagaagat 420
agtgcgtgac acctgcagga acatcggtt cgtctcaaat gatgtgggac tgggatgccg 480
acaactgcaa ngtcct 496

<210> 1905
<211> 247
<212> nucleic acid
<213> Glycine max
<400> 1905

nannatcgcn tgcacggta cgtnacgtcg gaattcggct cgagnntgga ttcattctctg 60
atgatgttgg tcttgatgct gacaaatgca aggtgttggc caacattgag cagcagagcc 120
ctgatatgcg ccagggtgtg cagggtcact tcaccaagcg cccagaggag gttggtgctg 180
gtgaccaggg tcacatgttt ggctatgcca ctgatgaaac ccctgagtac atgcccctca 240
gccatnt 247

<210> 1906
<211> 308
<212> nucleic acid
<213> Glycine max
<400> 1906

atncacatgt cacangcacg cgtacgtaag ctcggaattc ggctcgagnc tntcncttgt 60
ntgtgnatgc tgacacttat ggactggga agatccctga caaggaaatc ctcagnattg 120
tgaaggagag ttttgacttc aggcctggnn tgatctccat caaccttgnt ctcaagaggg 180
gtggaaatgg caggttcttg aagactgctg catatggnc ctttggcaga natgaccng 240
acttcacatg ggaantggtn angcgactca aggggganna ggtaccagct tancatanaag 300
ggntcct 308

<210> 1907
<211> 292
<212> nucleic acid
<213> Glycine max

<400> 1907

ancgnactgc acgcgtacgt aagctcggaa ttccggctcga gaaagtataa agatggcaga 60
gncattccta tttaacctcag agtcgggtgaa cgaggggacac cctgacaagc tctgcgacca 120
aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcttg 180
cgaaacatgc accaaaacca acttgggtcat ggtcttcgga gaaatcacga ccaaggccaa 240
cgttgactac gagaagatat ngngtgacac ctgcaggaac atcggtctcg tc 292

<210> 1908

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 1908

agtcgcangc acgcgtacgt aanctcggaa ttccggctcga ggtttggtta tgccactgta 60
tgaaaccctt gactacatgc cctcagcca tgccttgca accaaactcg gtgctgcct 120
caccgaggtt aggaataatg gtacctgtgc ttggctgagg ccagatggca agacacnagt 180
aactgttgag tactacaatg acaatgggtgc catgggtcca gttcgtgtcc aactgtcct 240
aatntncacc caacatgatg annngtgag caatgaccaa attgctgctg gaccttaaag 300

<210> 1909

<211> 458

<212> nucleic acid

<213> Glycine max

<400> 1909

atttcaggtg gtcttatagg ccaanaatga cgtaagacgc acgcgtncgt aanctcggaa 60
ttccggctcga gggccanggc aagccccact caaccaccac acctctctc gttcacgcta 120
ccctttctg ctcttcttct acctttcaag ttttaaaagt ataaagatgg cagagacatt 180
cctatttacc tcagagtcgg tgaacgaggg acaccctgac aagctctgcg accaaatctc 240
cgatgctgtc ctgcagcgtt gcctcgagca ggaccanac agcaaagttg cctgcgaaac 300
atgcacaaa accaacttgg ncatggtctt cgagaaaatc acgaccaagg ccaacgttga 360
ctacgagaag atagtgcgtg acacctgcag gaacatcggg ctctgtctca aatgatgtgg 420
gactggatgc cgacaantgc aangttctcg tcaacant 458

<210> 1910
 <211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 1910

 ngncncangc ncgcntacgt nanctcggaa ttcggtctga gctaccatct tctctctctc 60
 ttnccttagtg cctccttgcc agangtnaaa atggcccaag aaacttncct atncacatct 120
 gantcagtga acgaggggca ccctgacaag ctctgtgacc agatctccga tgctgtgctc 180
 gntgcatgct tggagcagga ccctgacagc aagggttgct gtgaaacctg caccaagacc 240
 aacatggtga tggtttttcgg agagatcaca accaaggcca acgtggacta tgagaagatt 300
 gtgcgtga 308

1000
 900
 800
 700
 600
 500
 400
 300
 200
 100
 0

<210> 1911
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 1911

 cgtgtacgta agctcggaa tctggctcgag aagtttttga agtatagaga tggcagagac 60
 attcctatatt acctcagagt cagtgaacga gggacaccct gacaagctct gtgaccaaatt 120
 ctctgatgct gtctctgacg cttgctctga acaggaccca gacagcaagg ttgcctgcca 180
 aacatgcacc aaaaccaact tggatcatggt cttcggagaa atcacgacca aggccaatgt 240
 ngactacgag aagatagtgc gtgacactgc aggacatngg tttgtccnaa cgnngngncn 300
 gncccc 306

<210> 1912
 <211> 504
 <212> nucleic acid
 <213> Glycine max

 <400> 1912

 aatgncaaac tncncagccg nctgncnng gtcataagggn ccgacngacc cgteccnaacc 60
 acgnatccgc tgnacantgn gcngacgcgt gggctgcnag aagacgacag aagggggcag 120
 cgcttgattt gaggccaggc aagccccact caaccaccac acctctctctc gttcacgcta 180

cccctttctg ctctttcttct acctttcaag ttttaaaagt ataaagatgg cagagacatt 240
 cctattttacc tcagagtcgg tgaacgaggg acaccctgac aagctctgcg accaaatctc 300
 cgatgctgtc ctcgacgctt gcctcgagca ggaccagac agcaaagttg cctgcgaaac 360
 atgcacaaaa accaacttgg tcatggtctt cggagaaatc acgaccaang ccaacgttga 420
 ctacgagaag atagtgentg acacctgcac ggaaatnggg ttcttctcaa ttaatttggg 480
 acgggtttcc cnaaaactnc aagg 504

<210> 1913
 <211> 289
 <212> nucleic acid
 <213> Glycine max

<400> 1913

tcnatgcacg cgtacgtaag ctcggaattc ggctcgagtc aaggtgtgca cggccacctt 60
 cacaagagg cctgaggaga ttggtgctgg tgaccaaggt catatgttcg gctatgccac 120
 tgacgagact cccgagctca tgcccttgag ccatgtcctt gccacgaagc cggtgccaag 180
 ctcaccgagg ttcggaagaa cgggacatgc ccttggttga gacctgatgg caagacccaa 240
 gtcactgttg agtactacaa tgacaagggt gccatggttc caatccgcg 289

<210> 1914
 <211> 345
 <212> nucleic acid
 <213> Glycine max

<400> 1914

gnattgnagt acgcgtacgt nagctcgga ttcggctcgn ggacttaaca acaacncana 60
 gngggttann gtctgttcaa gctaccatct ctctctctct ttcttagtgn ctctttgcna 120
 gaagttaana tggccaaga nactttccta ttcacatctg aatcagtga cgagggggcac 180
 cctgacaagc tctgtgacca natctccgat gctgtgctcg atgcatnctt ggagcaggac 240
 cctgacagca aggttgcttg tgaaacctgc accaanacna acatggtgat tgttttcgga 300
 gagatcacia ccaaggccaa cgttgactat gagaagattg tgcnt 345

<210> 1915
 <211> 331
 <212> nucleic acid

<213> Glycine max

<400> 1915

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60
ggttactgtc tgttcaagct accatctctc tctctcttct ntagtgctc cttgccagaa 120
gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacaacca aggccaacgt ggactatgag a 331

<210> 1916

<211> 244

<212> nucleic acid

<213> Glycine max

<400> 1916

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc caaggccaac gtagactatg 60
aaaagattgt ccgcgacaca tgccgcgaaa ttggattcat ctctgatgat gttgggtcttg 120
atgctgacaa atgcaagggtg ttggtcaaca ttgagcaaca gagcccggat atgcccagg 180
gtgtgcacgg ccacttcacc aagcgcccag aggaggttgg tgctgggtgac cagggtcaacn 240
tggt 244

<210> 1917

<211> 290

<212> nucleic acid

<213> Glycine max

<400> 1917

nnngtngcat gcncgcgtac gtaagctcng nntncggctc tntgcctata tcgtgagnca 60
ggctgcnaag agtggtgtgg naaatggcct tgccagaagg tgcattgtcc nagtttctta 120
tnccattggt gtccttgagc ccttgctcagt gtttatggac acttatggaa ctgggaanat 180
tcctgacaag gngattcttc aaattgtgna ggagaatttc gacttcagac ctggaatgat 240
caccattaac ttggacctta agaggggtgg ccataggttc ctcaagacag 290

<210> 1918

<211> 314
 <212> nucleic acid
 <213> Glycine max

 <400> 1918

 nnaagttnan gcacgcntac gtaagctcgg aattcggctc gaggttactg tctgttcaag 60
 ctaccatctc tctctctctt tottagtgcc tcttgccag aagttaaaat ggccaagaa 120
 actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag 180
 atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa ggttgctctg 240
 gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaa caaggccaac 300
 gtggactatg agaa 314

<210> 1919
 <211> 311
 <212> nucleic acid
 <213> Glycine max

 <400> 1919

 tattcnnnac gtgcgatgca cgcgtacgta agctcggaat tcggctcgag cactctctct 60
 gctottcttc tacctttcaa gtttttaaag tattaagatg gcagagacat tcctatttac 120
 ctcagagtca gtgaacgagg gacaccctga caagctctgc gaccaaactc ccgatgctgt 180
 octcgacgct tgccttgaac aggaccaga cagcaagggt gcctgcgaaa catgcaccaa 240
 gaccaacttg gtcattggtc tcggagagat caccaccaag gccaacgttg actacgagaa 300
 gatcgtgcgt g 311

<210> 1920
 <211> 281
 <212> nucleic acid
 <213> Glycine max

 <400> 1920

 ngcangcacg cgtacgtaag ctcggaattc ggctcgaggc ctcgagcagg acccagacag 60
 caaagttgcc tgcgaaacat gcacaaaac caacttggtc atggtcttcg gagaaatcac 120
 gaccaaggcc aacgttgact acgagnagat agtgcgtgac acctgcagga acatcggtt 180
 cgtctcaa at gatgtgggac tggatgccga caactgcaag gtctctgtca acattgagca 240

gcaganccct gatattgctc aggggtgttnc ccggccacct t 281

<210> 1921
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 1921

tccgaagtct cangcacgcy tacgtaagct cgggaattcgg ctcgagntac ctatgggtggg 60
 tgggggtgctc atgggtggagg tgccttttca ggggaaggacc ctaccaaggt tgacagaagt 120
 ggtgcctata tcgtgaggca ggctgcaaag agtggtgtgg caaatggcct tgcnaaaang 180
 gtgcnntggc cnangttttn aaggccatng gtgtccctga gcccttgtca gtgtttgtgg 240
 acacttatgg aactgggaag attcctgaca aggnngttct tcaaattgtg aaggngantt 300
 cngncttcng acntg 315

<210> 1922
 <211> 259
 <212> nucleic acid
 <213> Glycine max
 <400> 1922

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ggtggagggtg ccttttcagg 60
 gaaggaccct accaaggttg acagaagtgg tgcctatata gtgaggcagg ctgcaaagag 120
 tgttgtggca aatggccttg cagaaggtgc attgtccaag tttcctatgc cattgggtgtc 180
 cctgagccct tgtcagtgtt tgnccgacact tatggaactg ggaagattcc tgacaaggag 240
 attcttcaaa ttgtgaagg 259

<210> 1923
 <211> 300
 <212> nucleic acid
 <213> Glycine max
 <400> 1923

agtcgcatgc acgcgtacgt aagctcggga attcggctcg agacctctca agtttttgaa 60
 gtatagagat ggcagngaca ttcctattta cctcagagtc agtgaacgag ggacaccctg 120
 acaagctctg tgaccaaatc tcatgtgtgc ctcgacgctt gcctogaaca ggacccagac 180

agcaaggttg cctgcgaaac atgcacaaaa accaacttgg tcatggtctt cggagaaatc 240
acgaccaagg ccaatgttga ctacgagaag atagtgcgtg acacctgcag gaacatcggt 300

<210> 1924
<211> 290
<212> nucleic acid
<213> Glycine max

<400> 1924

anncgcangc acgcgtacgt aagctcggaa ttcggtctga gtcaagtttn tgaagtatag 60
agatggcaga gacattccta ttacctcag agtcagtga cgagggacac cctgacaagc 120
tctgtgacca aatctctgat gctgtcctcg acgcttgctt cgaacaggac ccagacagca 180
aggttgcctg cgaaacatgc accaaaacca acttgggtcca tggctcttcgg agaaatcacg 240
accaaggcca atgttgacta cgagaagata gtgcgtgaca cctgcaggaa 290

<210> 1925
<211> 294
<212> nucleic acid
<213> Glycine max

<400> 1925

ngtcgcatgc acgcgtacgt aagctcggaa ttcggtctga gcggtctgag cggtctgagg 60
gcaaattggcc ttgccagaag gtgcattgtc caagtttctt atgccattgg tgtcnctgag 120
ccottgtcag tgtttgtgga cacttatgga actgggaaga ttcttgacaa ggagattctt 180
caaattgtga aggagaattt cgacttcaga cctggaatga tcaccattaa cttggacctt 240
aagaggggtg gccataggtt cctcaagaca gctgcttatg gacactttgg aagg 294

<210> 1926
<211> 473
<212> nucleic acid
<213> Glycine max

<400> 1926

anctttgtac gcgccccagg taccggtaaa ggaattccng gctcgaccca cgcgtaagcc 60
cacgcgtccg tacggctgcg agaagacgac agaagggggc agcgcttgat ttgaggccag 120
gcaagcccca ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt 180

ctacctttca agtttttaaaa gtataaagat ggcagagaca ttcctattta cctcagagtc 240
 ggtgaacgag ggacaccctg acgagctctg cgaccaaadc tccgatgctg tcctcgacgc 300
 ttgcctcgag caggacccag acagcaaagt tgccctgcgaa acatgcacca aaaccaactt 360
 ggtcatggtc ttccggagaaa tcacgaccaa ggccaacgtt gactacgaag aagatagtg 420
 gtgacacctg cangaacatc ggcttcgtct caaatgatgt tggaactgga tgc 473

<210> 1927
 <211> 490
 <212> nucleic acid
 <213> Glycine max

<400> 1927

Sequence 1

atttnacctt cccnggccn gttaaaggta aaganttccc gggncgaccc acgcgtccgc 60
 ccacgcgtcc gtacggctgc gagaagacga cagaaggggg gcagcgcttg atttgaggcc 120
 aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt totgctcttc 180
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 240
 tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgcctcgac 300
 gcttgccctg agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaac 360
 ttggtcatgg tcttcggaga aatcacgacc aaggccaacg ttgactacga agaagatagt 420
 gcgtgacacc tgcaggaaca tcggcttcgt ctcaaaatga tgtgggactg gatgccgaca 480
 actgnnangg 490

<210> 1928
 <211> 320
 <212> nucleic acid
 <213> Glycine max

<400> 1928

aanagtcgca ngcacgctta cgtnaagctc ggaattcggc tcgagaaagc gggntactgt 60
 ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga agttaaaatg 120
 gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag 240
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300

aaggccaacg tggactatgn

320

<210> 1929
<211> 294
<212> nucleic acid
<213> Glycine max

<400> 1929

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ggagattctc aacattgtga 60
aggaaaactt tgatttcagg cctgggatga tctccatcaa ccttgatctc aagaggggtg 120
gaaataacag gtttttgaag actgctgcct atggacactt tggaagagaa gaccctgact 180
tcacatggga agtgggtcaaa cccctcaagt gggagaaggc ctaagtaatt cattccactg 240
ctctatgctg gaagtttttt gagcggtgcc cttataatat gtctaataatc catn 294

<210> 1930
<211> 304
<212> nucleic acid
<213> Glycine max

<400> 1930

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ggagattctc aacattgtga 60
aggaaaactt tgatttcagg cctgggatga tctccatcaa ccttgatctc aagaggggtg 120
gaaataacag gtttttgaag actgctgcct atggacactt tggaagagaa gaccctgact 180
tcacatggga agtgggtcaaa cccctcaagt gggagaaggc ctaagtaatt cattccactg 240
ctctatgctg gaagtttttt gagcggtgcc cttataatat gtctaataatc cataactttc 300
cacg 304

<210> 1931
<211> 321
<212> nucleic acid
<213> Glycine max

<400> 1931

cgcacatgcacg cgtacgtnag ctcggaattc ggctcgagct tctacctctc aagtttttga 60
agtatagacn ncggcagaga cattccctat ttaccttcag agttcagtga acgagggaca 120
cnctgacaag ctctgtgacc aaatctctga tgctgtcctc gacggttgcc tcgaacagga 180

cccagacagc naggttgcct gcgaaacatg caccaaaacc aacttgggtca tgggtcttcgg 240
 agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca cctgcaggaa 300
 catcggcttt gtctcaaacg a 321

<210> 1932
 <211> 281
 <212> nucleic acid
 <213> Glycine max

<400> 1932

cgcatgcacg cgtacgtaag ctoggaattc ggctcgagct tctacctctc aagtttttga 60
 agtatagaga tggcagagac attcctatctt acctcagagt cagtgaacga gggacacctt 120
 gacaagctct gtgaccaaatt ctctgatgct gtctctgacg cttgcctcga acaggaccca 180
 gagagcaagg ttgcctgcga aacatgcacc aaaaccaact tgggtcatggc cttcggagaa 240
 atcacgacca aggccaatgt tgactacgag aaganagtgc g 281

<210> 1933
 <211> 292
 <212> nucleic acid
 <213> Glycine max

<400> 1933

natacatgca cgcgtacgta agctcggaat tcggctcgag ctctctgctc ttcttctacc 60
 tttcaagttt ttaaagtatt aagatggcag agacattcct atttacctca gagtcagtga 120
 acgagggaca cctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 180
 ttgaacagga cccagacagc aaggttgcct gcgaaacatg caccaagacc aacttgggtca 240
 tgggtcttcgg agagatcacc accaaggcca acgttgacta cgagaagatc gt 292

<210> 1934
 <211> 266
 <212> nucleic acid
 <213> Glycine max

<400> 1934

ctctgcgncc aaatctccga tgctgtcctc gacgcttgcc ttgaacagga cccagacagc 60
 aaggttgcct gcgaaacatg caccaagacc aacttgggtca tgggtcttcgg agagatcacc 120

accaaggcca acgttgacta cgagaagatc gtgcgtgaca cctgnaggaa catcggttc 180
 gtctcaancg atgtgggact tgatgctgac aactgccaaag gtnctgnaa acattgaggn 240
 nncagagccc tggatattgc ccaggg 266

<210> 1935
 <211> 310
 <212> nucleic acid
 <213> Glycine max

<400> 1935

cgcnaaangc gtacgtnagc tcggaattcg gctcgagncg ggttactgtc tgttcaagct 60
 accatctctc tctctctttc ttagtgctc cttgccagaa gttaaaatgg cccaagaaac 120
 tttcctattc acatctgaat cagtgaacga ggggcaccct gacaagctct gtgaccagat 180
 ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg ttgcctgtga 240
 aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca aggccaacgt 300
 ggactatgag 310

<210> 1936
 <211> 299
 <212> nucleic acid
 <213> Glycine max

<400> 1936

gtgcgancga cgcgtacgta agctcggaat tcggctcgag gcaacagagt cctgatattg 60
 ctcaagggtgt gcacggccac ctcaaaaga ggctgagga gattggtgct ggtgaccaag 120
 gtcatatgtt cggctatgcc actgacgaga ctcccgagct catgcccttg agccatgtcc 180
 ttgccacgaa gctcgggtgcc aagctcaccg aggttcggaa gaacgggaca tgcccttggc 240
 tgagacctga tggaccactg ntgantgatt acgatcacga ttaattcggc cccgacagt 299

<210> 1937
 <211> 311
 <212> nucleic acid
 <213> Glycine max

<400> 1937

ncnacgcan gcacgcgtac gtnagctcgg aattcggctc gagctctcaa gtttttgaag 60

tatagagatg gcagagacat tcctattttac ctcagagtca gtgaacgagg gacaccctga 120
 caagcttctg tgaccnaaat ctctgatgct gtcctcgacg cttgcctcga acaggaccca 180
 gagagcaagg ttgcctgcga aacatgcacc aaaaccaact tggatcatggt cttcggagaa 240
 atcacgacca aggccaatgt tgactacgag aagatagtgc gtgacacctg caggaacatc 300
 ggctttgtct t 311

<210> 1938
 <211> 319
 <212> nucleic acid
 <213> Glycine max

<400> 1938

gatgcacgcg tacgtaagct cggaattcgg ctcgagcaca aagcgggtta ctgtctgttc 60
 aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa aatggcccaa 120
 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caangttgcc 240
 tgtgaaacct gcaccaagac caacatggta tggttttcgg agagatcaca accaaggcca 300
 acgtggacta tgagaagat 319

<210> 1939
 <211> 315
 <212> nucleic acid
 <213> Glycine max

<400> 1939

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cggctcgagg cacgctctgc 60
 ttccagcgag tgttttttct tcgtttcaac accttaattt gcanacgctg cttcttcccg 120
 cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc 180
 ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcott gaacaggacc 240
 ctgacagcaa ggttgctgtg gagacatgca ccaagaccaa catggtcatt gtctttggag 300
 agatcacaac caagg 315

<210> 1940
 <211> 303
 <212> nucleic acid

<213> Glycine max

<400> 1940

cgcangcacg cgtacgtaag ctcggaattc ggctcgaggt cttgatgctg acaactgcaa 60
ggtcctcgtc aacattgagc aacagagtcc tgatattgct caaggtgtgc acggccacct 120
cacaagaggg cctgaggaga ttggtgctgn tgaccaaggt catatgttcg gctatgccan 180
tganganact cccgagctca tgccttgag ccatgtcctt gccacgaagc tcggtgccaa 240
gtctcaccga ggctnggnag aacgggacat ccctgggnt gagacntgnt ggcaaagncc 300
aaa 303

<210> 1941

<211> 335

<212> nucleic acid

<213> Glycine max

<400> 1941

tcgtgcacg cgtacgtaag ctcggaattc ngctcgaggc ccactcaac caccacacct 60
ctcctcgctt acgtacccc ttctgtctct tcttctacct ttcaagtttt aaaagtataa 120
agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc 180
tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca 240
aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaaatcacga 300
ccaaggccaa cgttgactac gagaagatag tgcgt 335

<210> 1942

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1942

tcgcangcnc gcgtacgtna gctcggaatt cggctcgagg ggattctcaa cattgtgaag 60
gaaaactttg atttcaggcc tggtatgatc tccatcaacc ttgatctcaa gaggggtgga 120
aataacaggt ttttgaagac tgctgcctat ggacactttg gaagagaaga cctgacttc 180
acatgggaag tggtcaaacc cctcaagtgg gagaaggcct aagtaattca ttccactgct 240
ctatgctgga agttttttga gcgttgccct ataatatgtc taata 285

<210> 1943
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 1943

 ngctctcangc acncgtacgt aagctnggaa ttccggctcna ggtctgttcn ngctaccatc 60
 tctctnctct ctttcttagt gcttccttgc cagaagttaa antggcccaa gaaactttcc 120
 tattcacatc tgaatnagtg aacgaggggc accctgacaa gctctgtgac cagatctccg 180
 atgctgtgct cgatgcatgc ttggagcagg accctgacag canngttgcc tgtgaaacct 240
 gcaccaagac caacatgggtg atggtttttcg gagagatcac naccaaggnc aacgtggact 300
 atgagaagat 310

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<210> 1944
 <211> 317
 <212> nucleic acid
 <213> Glycine max

 <400> 1944

 gtgcgatgca cgcgtacgta agctcgggaa ttccggctcga ggtaggttc tgcacgctct 60
 gcttccagcg agtggttcttt cttcgtttca acaccttaat ttgcacacgc tgctttctca 120
 gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa cgaggggtcac 180
 cccgacaagc tgtgcgacca gatctctgat gcagtgtcgc atgcgtgcct tgaacaggac 240
 cctgacagca aggttgacctg tgagacatgc accaagacca acatgggtcat ggtcttttga 300
 gagatcacia ccaaggc 317

<210> 1945
 <211> 331
 <212> nucleic acid
 <213> Glycine max

 <400> 1945

 tngnnngcac gcgtacgtaa gctcgggaatt cggctcgagc ggctcgagtt tgggagttag 60
 gttctgcacg ctctgcttcc agcgagtgtt ctttcttcgt ttcaacacct taatttgcac 120
 acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180

taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt 240
gccttgaaca ggaccctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg 300
tcattggtctt tggagagatc acaaccaagg c 331

<210> 1946
<211> 314
<212> nucleic acid
<213> Glycine max

<400> 1946

nncacgcgta cgtaagctcg gaattcggct cgagnggagt taggttctgc acgctctgct 60
tccagcgagt gttctttctt cgtttcaaca ccttaatttg cacacgctgc ttcttcngct 120
tgagaaatgg cacaagaaac ctttctatct acatctgaat ctgtaaacga gggtcacccc 180
gacaagctgt ggcaccagat ctctgatgca gtgctcgatg cgtgccttga acaggaccct 240
gacagcaagg ttgcctgtga gacatgcacc aagaccaaca tggatcatggc ctttggagag 300
atcacaacca aggc 314

<210> 1947
<211> 306
<212> nucleic acid
<213> Glycine max

<400> 1947

ctcgnatgca cgcgtacgta agctcggaat tcggctcgag gcacgctctg cttccagcga 60
gtgtttctttc ttcgtttcaa caccttaatt tgcacacgct gcttcttcag cttgagaaat 120
ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc ccgacaagct 180
gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt gaacaggacc ctgacagcaa 240
ggttgcctgt gagacatgca ccaagaccaa catggatcatg gtctttggag agatcacaac 300
caaggc 306

<210> 1948
<211> 297
<212> nucleic acid
<213> Glycine max

<400> 1948

atngnagtcg cangcncgcg tacgt nagct cggaattcgg ctcgaggnga ttctcaacat 60
 tgtgaaggaa aactttgatt tcaggcctgg tatgatctcc atcaaccttg atctcaagag 120
 ggggtggaaat aacaggtttt tgaagactgc tgcctatgga cactttggaa gagaagaccc 180
 tgacttcaca tgggaagtgg tcaaaccct caagtgggag aaggcctaag taattcattc 240
 cactgctcta tgctggaagt tttttgagcg ttgcccttat aatatgtcta atatcca 297

<210> 1949
 <211> 217
 <212> nucleic acid
 <213> Glycine max

<400> 1949

gcgtacgtaa gctcggaatt cggctcgagg acaaatgcaa ggtgttggtc aacattgagc 60
 agcagagccc tgatatcgcc caggggtgtgc acggtcactt caccaagcgc ccagaggagg 120
 ttggtgctgg tgaccaggtt cacatgtttg gctatgccac tgatgaaacc cctgagtaca 180
 tgccctcag ccatgtcctt gcaaccaaac tcgggtgc 217

<210> 1950
 <211> 291
 <212> nucleic acid
 <213> Glycine max

<400> 1950

agnnaangca cgcgtacgta agctcggaat tcggctcgag actctctctg ctcttcttct 60
 acctttcaag tttttaaggt attaagatgg cagagacatt cctatttacc tcagagtcag 120
 tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctgacgctt 180
 gcottgaaca ggaccagac agcaagggtg cctgcgaaac atgcaccaag accaacttgg 240
 tcatggtctt cggagagatc accaccaagg ccaacgttga ctacgagaag a 291

<210> 1951
 <211> 291
 <212> nucleic acid
 <213> Glycine max

<400> 1951

gtcgcanntt ngcgtacgta agctcggaat tcggctcgag actctctctg ctcttcttct 60

acctttcaag tttttaaggt attaagatgg cagagacatt cctatattacc tcagagtcag 120
 tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt 180
 gccttgaaca ggaccagac agcaaggttg cctgcgaaac atgcaccaag accaacttgg 240
 tcatggtctt cggagagatc accaccaagg ccaacgttga ctacgagaag a 291

<210> 1952
 <211> 319
 <212> nucleic acid
 <213> Glycine max
 <400> 1952

gtgcgatgca cgcgtacgta agctcggaat tcggctcgag gttaggttct gcacgctctg 60
 cttccagcga gtgttctttc ttcgtttcaa caccttaatt tgcacacgct gcttcttcag 120
 cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc 180
 ccgacaagct gtgcgaccag atctctgatg cagtgtctga tgcgtgcctt gaacaggacc 240
 ctgacagcaa ggttgctgtg gagacatgca ccaagaccaa catggtcatg gtctttggag 300
 agatcacaac caaggccag 319

<210> 1953
 <211> 288
 <212> nucleic acid
 <213> Glycine max
 <400> 1953

gtcgnangca cgcgtacgta agctcggaat tcggctcgag tctctctgtt ctcttctacc 60
 tctcaagttt ttgaagtata gagatggcag agacattcct atttacctca gagtcagtga 120
 acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 180
 tcgaacagga ccagacagc aaggttgctt gcgaaacatg caccaaaacc aacttggtca 240
 tgggtcttcgg agaaatcacg accaaggcca atgttgacta cgagaaga 288

<210> 1954
 <211> 248
 <212> nucleic acid
 <213> Glycine max
 <400> 1954

acctctcaag tttttgaagt atagagatgg cagagacatt cctattttacc tcagagtcag 60
 tgaacgaggg acaccctgac aagctctgtg accaaatctc tgatgctgtc ctcgacgctt 120
 gcctcgaaca ggaccagac agcaagggtg cctgcgaaac atgcaccaaa accaacttgg 180
 tcatggtctt cggagaaatc acgaccaagg ccaatgttga ctacgagaag atatgctga 240
 cactgcag 248

<210> 1955
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 1955

cgcangcacg cgtacgttag ctcggaattc ggctcgagnt ttgaaccctt ctggcaggtt 60
 tgtcattgga gggccgcatg gngatgctgg tctcaccggc ngcaagntca ncatcgacac 120
 ctatgntcng atgggggtgca catggtggtg gtgccttctn tgggaaggat ccgccaangt 180
 tgataggagt ggtgcctaca ttgtgaggca agctgcaaag agcattgttn caaatggant 240
 tgctaggagg gcaattgtgc aagtttcta tgccattggt gtgcctganc cntgtctgtg 300
 nttgttnac 309

<210> 1956
 <211> 292
 <212> nucleic acid
 <213> Glycine max
 <400> 1956

cgtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gggaggaggt gccttttcag 60
 ggaaggaccc taccaagggt gacagaagtg gtgcctatat cgtgaggcag gctgcaaaga 120
 gtgttggtggc aaatggcctt gccagaaggt gcattgtcca agtttctat gccattggtg 180
 tccctgagcc ctgtcagtgt ttgtggacac ttatggaact gggaagattc ctgacaagga 240
 gattcttcaa atgtgaagga gaattcgact tcagacctgg aatgatcacc at 292

<210> 1957
 <211> 317
 <212> nucleic acid
 <213> Glycine max

<400> 1957

tgcatgcacg cgtacgtaag ctcggaattc ggctcgagta acaacagcac aaagcggggt 60
actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg ccagaagtta 120
aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctgaca 180
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240
gcaagggttg ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 300
caaccaaggc caacgtg 317

<210> 1958

<211> 219

<212> nucleic acid

<213> Glycine max

<400> 1958

tagtgectcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgottggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
ggatgatggt ttcggagaga tcacaaccaa ggccaacgt 219

<210> 1959

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 1959

togatgcaen cgtacgtgag ctcggaattc ggctcgaggc gggtnnctgt ctgttcaagc 60
taccatctct ctctctcttt cttagtgcct ccttgccaga agttaaagt gccaagaaa 120
ctttcttatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc tgtgaccaga 180
tctccgatgc tgtgctcgat gcatgcttg agcaggaccc tgacagcaag gttgcctgtg 240
aaacctgcac caagaccaac atggtgatgg ttttcggaga ggtcacaacc aaggccaacg 300
tgatatgan aagattgtgc gtgac 325

<210> 1960

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 1960

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcacaaagcg ggttactgtc 60
tggtcaagct accatctctc tctctcttct ttagtgcttc cttgccagaa gttaaaatgg 120
cccaagaaac ttctctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180
gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240
ttgcctgtga aacctgcacc aagaccaaca tggatgatgt ttcggagag atcacaacca 300
aggccaacgt ggactn 316

<210> 1961

<211> 495

<212> nucleic acid

<213> Glycine max

<400> 1961

ggnnnnnnnn aatttactct gcccgncagg tacangtaca gaattcccgg ntcgaccac 60
gcgtcagtac ggctgcgaga agacgacaga agggggcagc gcttgatttg aggccaggca 120
agccccactc aaccaccaca cctctctctg ttcacgctac ccctttctgc tcttcttcta 180
ccctttcaagt tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt 240
gaacgaggga caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg 300
cctcgagcag gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttgg 360
catggtcttc ggagaaatca cgaccaangc caacgttgac tacganaaga tatgcgtgac 420
acctgcaagg aacatcggtt tctctcaaat gatgttgga ctggatgccg acaactgcaa 480
ggtctcgtca acatt 495

<210> 1962

<211> 270

<212> nucleic acid

<213> Glycine max

<400> 1962

agtcgngca cgcgtacgta aactcggaat tcggctcnag cnaggttgat aggnatggtg 60
cttacattgt gagacaggct gctaaganca ttgtggcaag tggacttgcc agaaggtgca 120

ttgtagcaag tgtcttange cattggtgtg cctgagcctt tgtctgtgtt tnttgacacc 180
tatggcactg ggaagatcca tgataaggag attctcaaca ttgngaagga aaactttgat 240
ttcangcctg gnatgatctc catcaacctt 270

<210> 1963
<211> 282
<212> nucleic acid
<213> Glycine max
<400> 1963

tgcaacaaaa ccaacttggg catggtcttc ggagaaatca cgancaaggc caacgnttga 60
ctaaggagaa gnnatgcgnt gaacacctgg caggnccatc ggcttcgtct caaatnangt 120
gggactgccc tgccgacaac tgcaagggtc tcgtcaacat tgagcagcag agccctgata 180
ttgctcaggg tgtacacggc caccttacca aaaaacctga agaaattggg gctgggtgacc 240
agggtcacat gtttggcnat gccactgatg aaaccctga ct 282

<210> 1964
<211> 306
<212> nucleic acid
<213> Glycine max
<400> 1964

nngttontgc acgcgtacgt cagctcggaa ttcggctcga ggcggttac tgtctgttca 60
agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaa atggcccaag 120
aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240
gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagnhnaca accaaggcca 300
acgtgg 306

<210> 1965
<211> 317
<212> nucleic acid
<213> Glycine max
<400> 1965

gcangcacgc gtacgtaagc tcggaattcg gctcgaggac ttaacaacag cacaagcgg 60

gttactgtct gttcaagcta ccactctctct ctctctttct tagtgctcc ttgccagaag 120
 ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcacctg 180
 acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggacctg 240
 acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttccgagaga 300
 ncacaaccaa ggccaag 317

<210> 1966
 <211> 296
 <212> nucleic acid
 <213> Glycine max
 <400> 1966

togtangtaa gctcggaatt cggctcgagg cgggncactg tctgttcaag ctaccatctc 60
 tctctctctt tcttagtgcc tccttgccag aagttaaaat ggcccaagaa actttcctat 120
 tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag atctccgacn 180
 ctgtgctoga tgcattgctg gancaggacc ctgacagcaa ggttgctgt gaaacctgca 240
 ccaagaccaa catggtgatg gttttcggag agatcacaac caaggccaac gtggat 296

<210> 1967
 <211> 318
 <212> nucleic acid
 <213> Glycine max
 <400> 1967

acgtcgcatg ctagegtacg taagctcgga attcggctcg agcacaagc gggttactgt 60
 ctgttcaagc taccatctct ctctctcttt cttagtgctt ccttgccaga agttaaagt 120
 gccaagaaa ctttctatt cacatctgaa tcagtgaacg aggggcacc tgacaagctc 180
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttg agcaggacc tgacagcaag 240
 gttgctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300
 aaggccacgt ggactatg 318

<210> 1968
 <211> 313
 <212> nucleic acid
 <213> Glycine max

<400> 1968

gtcncatgca cgcgtacgta agctcggaat tcggctcgag acagcacaaa gcgggttact 60
gtntgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120
tgggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180
ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240
aaggttgccct gtgaaacctg caccaagacc aacatggtga tggttttcgg agagatcaca 300
accaaggcca acg 313

<210> 1969

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 1969

ncgtcgcatg cgcgtacg taagctcgga attcggtcgc agngcagttt taaaagtata 60
aagatggcag agacattcct atttacctac agagtcggtg aacgagggac accctgacaa 120
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 180
caaagttgcc tgcgaaacat gcacaaaaac caacttggtc atggttttcg gagaaatcac 240
gaccaaggcc aacgttgact acgagaagat agtgcgtgac acctgcagga a 291

<210> 1970

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 1970

ntgcantnac gcgtacgtaa gctcggaatt cggctcnagn cagacttaac aacagcacia 60
agcgggttac tgtctgttca agctaccatc tctctcncct ctttcttagt gcctccttgc 120
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
accctgacag caaggttgcc tgtgaaacct gcancaagac caacatggtg atggtttttcg 300
gagagatcac aaccaaggcc aacgtgg 327

<210> 1971

<211> 294
 <212> nucleic acid
 <213> Glycine max

 <400> 1971

 tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc cacaccactc tctctgnntc 60
 ttctttctacc tttcaagtnt ttaaagtatt aagatggcag agacattcct atttacctca 120
 gagtcagtga acgagggaca ccttgacaag ctctgcgacc aaatctccga tgctgtcctc 180
 gacgcttgcc ttgaacagga ccagacagc aagggtgcct gcgaaacatg caccaagacc 240
 aacttggtca tgggtcttcgg agagatcacc accaaggcca acgttgacta cgag 294

<210> 1972
 <211> 293
 <212> nucleic acid
 <213> Glycine max

 <400> 1972

 gtgcgatgca cgcgtacgta agctcggaat tcggctcgag ntttcaagtt tttaaagtat 60
 taagatggca gagacattcc tatntacctc agagtcagtg aacgagggac accctgacaa 120
 gctctgcgac caaatctccg atgctgtcct cgacgcttgc cttgaacagg acccagacag 180
 caaggttgcc tgcgaaacat gcaccaagac caacttggtc atggtcttcg gagagatcac 240
 caccaaggcc aacgttgnet acgagaagtc gtgcgtgaca ctgaggaaca tcg 293

<210> 1973
 <211> 339
 <212> nucleic acid
 <213> Glycine max

 <400> 1973

 tcgngcacg cgtacgtaag ctcggaattc ggctcgagcg agccatttgg gagttagggtt 60
 ctgcacgctc tgcttcacgc gagtgttctt tcttcgnttc aacaccttaa tttgcacacg 120
 ctgcttcttc agcttgagaa atggcacaag aaacctttct attcacatct gaatctgtaa 180
 acgaggggtca ccccgacaag ctgtgcgacc agatctctga tgcagtgctc gatgcgtgcc 240
 ttgaacagga ccctggacag caaggttgcc tgtgagacat gcaccaagac caacatggtc 300
 atggtctttg gagagatcac aaccaaggcc aacgtagat 339

<400> 1974

[illegible]

<400> 1975

Year	Age	Sex	Location	Occupation	Education	Marital Status	Religion	Health Status	Family Size	Income	Assets	Debt	Life Expectancy	Mortality Rate	Healthcare Access	Life Quality	Life Satisfaction	Life Expectancy	Mortality Rate	Healthcare Access	Life Quality	Life Satisfaction
1990	20	Male	Urban	Student	High School	Single	Christian	Good	2	\$10,000	\$5,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
1995	25	Female	Rural	Farmer	High School	Married	Muslim	Fair	3	\$15,000	\$10,000	\$5,000	70	120	No	7.5	8.0	70	120	No	7.5	8.0
2000	30	Male	Urban	Teacher	College	Single	Christian	Good	2	\$20,000	\$15,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2005	35	Female	Rural	Nurse	College	Married	Muslim	Good	3	\$25,000	\$20,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2010	40	Male	Urban	Engineer	College	Single	Christian	Good	2	\$30,000	\$25,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2015	45	Female	Rural	Doctor	College	Married	Muslim	Good	3	\$35,000	\$30,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2020	50	Male	Urban	Lawyer	College	Single	Christian	Good	2	\$40,000	\$35,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2025	55	Female	Rural	Professor	College	Married	Muslim	Good	3	\$45,000	\$40,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2030	60	Male	Urban	Retired	College	Single	Christian	Good	2	\$50,000	\$45,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2035	65	Female	Rural	Retired	College	Married	Muslim	Good	3	\$55,000	\$50,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2040	70	Male	Urban	Retired	College	Single	Christian	Good	2	\$60,000	\$55,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2045	75	Female	Rural	Retired	College	Married	Muslim	Good	3	\$65,000	\$60,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2050	80	Male	Urban	Retired	College	Single	Christian	Good	2	\$70,000	\$65,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2055	85	Female	Rural	Retired	College	Married	Muslim	Good	3	\$75,000	\$70,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2060	90	Male	Urban	Retired	College	Single	Christian	Good	2	\$80,000	\$75,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2065	95	Female	Rural	Retired	College	Married	Muslim	Good	3	\$85,000	\$80,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2070	100	Male	Urban	Retired	College	Single	Christian	Good	2	\$90,000	\$85,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2075	105	Female	Rural	Retired	College	Married	Muslim	Good	3	\$95,000	\$90,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2080	110	Male	Urban	Retired	College	Single	Christian	Good	2	\$100,000	\$95,000	\$5,000	75	100	Yes	8.5	9.0	75	100	Yes	8.5	9.0
2085	115	Female	Rural	Retired	College	Married	Muslim															

<400> 1976

tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atggatgatg ttttcggaga 300
 gatcacaacc aaggc 315

<210> 1977
 <211> 316
 <212> nucleic acid
 <213> Glycine max

<400> 1977

gtgcgacga cgcgtagcta agctcggaat tcggctcgag gacttaacaa cagcacaaag 60
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120
 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggatgatg gttttcggag 300
 agatcacaac caaggc 316

<210> 1978
 <211> 309
 <212> nucleic acid
 <213> Glycine max

<400> 1978

nnagangcac gnacaacgta agctcggaat tcggctcgag caacagcaca aagcgggtta 60
 ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120
 aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180
 gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240
 caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggttttcg gagagatcac 300
 aaccaaggc 309

<210> 1979
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 1979

natgnntacg tnagctcgga attcggctcg agcagcacia agcgggttac tgtctgttca 60
 agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaaa atggcccaag 120
 aaacttttct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180
 agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240
 gtgaaacctg caccaagacc aacatggtga tggttttcgg agagatcaca accaaggc 298

<210> 1980
 <211> 314
 <212> nucleic acid
 <213> Glycine max

<400> 1980

togcangcac gcgtacgtaa nctcggaatt cggtctgagn ttgggagtta ggttctgcac 60
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgtgctt 120
 cttcagcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180
 gtcaccccgca caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccctgaac 240
 aggaccctga cagcaagggt gctgtgaga catgcaccaa gaccaacatg gtcattggtct 300
 ttggagagat caca 314

<210> 1981
 <211> 325
 <212> nucleic acid
 <213> Glycine max

<400> 1981

gtgcgatgca cgcgtacgta agctcggaat tcgggctcga gcttaacaac agcaciaaagc 60
 gggttactgt ctgttcaagc taccatntct ctctctcttt cttagtgcct ccttgccaga 120
 agttaaaatg gcccaagnaa acttttctat tcacatctga atcagtgaac gaggggcacc 180
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300
 agatcacaac caaggccaac gtggn 325

<210> 1982
 <211> 315
 <212> nucleic acid

<213> Glycine max

<400> 1982

gtcgcangca cgcgtacgta agctcggaat tcggctcgag nttgggagtt aggtttctgca 60
cgctctgctt ccagcgagtg ttctttcttc gtttcaacac cttaatttgc acacgctgct 120
tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180
ggtcaccccg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240
caggaccctg acagcaaggt tgctgtgag acatgcacca agaccaacat ggtcatggtc 300
tttgagagaga tcaca 315

<210> 1983

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 1983

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cctcagccat gtccttgcaa 60
ccaaacttgg tgctcgcttc acagagggtta ggaagaatgg cacctgtgct tggttgaggc 120
cagatggtaa gacacaagta accgtcgagt actacaatga caatgggtgcc atggttccag 180
ttcgtgtcca cactgtccta atttccaccc aacatgatgn ncctgtgagc aatgatcaaa 240
ttcgtgogga cttaaaggca tgttataaac ctgncatccn ggaaaatact tgnaggaa 298

<210> 1984

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 1984

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcgggttact gtctgttcaa 60
gtaccatct ctctctctct nnccttagtg ctccttgcc agaagttaa atggccaag 120
aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240
gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagatcaca accaaggcc 299

<210> 1985

<211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 1985

 ncgtnncntn nagctcggaa ttcggctcga gcttaacaac agcacaaagc gggttactgt 60
 ctgttcaagc taccatctct ctctctcttt cttagtgcct cottgccaga agttaaaatg 120
 gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag 240
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300
 aaggcc 306

<210> 1986
 <211> 300
 <212> nucleic acid
 <213> Glycine max

 <400> 1986

 gtencatgca cgcgtacgta agctcggnat tcggctcgag aagcgggtta ctgtctgttc 60
 aagetaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa aatggcccaa 120
 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caaggttgcc 240
 tgtgaaacct gcaccaagac caacatggtg atggttttcg gagagatcac aaccaaggcc 300

<210> 1987
 <211> 319
 <212> nucleic acid
 <213> Glycine max

 <400> 1987

 gaaccgtngc tggtagctaa gctcgggaatt cggctcgagg cagacttaac aacagcacia 60
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
 agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180
 cctgacaaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
 cctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300

agagatcaca accaaggcc 319

<210> 1988
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 1988

gnannngcac gcgtacgtaa gctcggaatt cggctcgaga acaacagcac aaagcgggtt 60
actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg ccagaagtta 120
aatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctgaca 180
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240
gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 300
caaccaaggc c 311

<210> 1989
<211> 331
<212> nucleic acid
<213> Glycine max

<400> 1989

ctngcangca gcgtacgtaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
agaagttaaa atgggcccac ganactttcc tntcacatc tgaatcagt aacgaggggc 180
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttcg 300
gagagatcac aaccaaggcc aacgtggact a 331

<210> 1990
<211> 319
<212> nucleic acid
<213> Glycine max

<400> 1990

tcaacagtcg catgcacgcg tacgtaagct cggaattcgg ctcgagaaca acagcacia 60
gcgggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctccttgcca 120

gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180
cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac 240
cctgacagca aggttgcttg tgaaacctgc accaagacca acatggtgat ggttttcgga 300
gagatcacia ccaaggccc 319

<210> 1991
<211> 288
<212> nucleic acid
<213> Glycine max
<400> 1991

ntaangcacg cgtacgtaag ctcggaattc ggctcgaggg acaaatgcaa ggtgttggtc 60
aacattgagc aacagagccc ggatatcgcc cagggtgtgc acggccactt caccaagcgc 120
ccagaggagg ttggtgctgg tgaccagggt cacatgtcac angtatgcc ncatgncac 180
ccccgagtac atgcccctca gccatgtcct tgcaacaaaa cttggtggnt cgccncacag 240
aggttaggag aattgcactg tgcttggttg aggccagatg gtaagaca 288

<210> 1992
<211> 333
<212> nucleic acid
<213> Glycine max
<400> 1992

nncnngcac gcgtacgtng ctcggaattc ggctcgagct taacaacagc acaaagcggg 60
ttactgtctg ttcaagctac catctctctc tctctttctt agtgcctcct tgccagaagt 120
taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg ggcaccctga 180
caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga 240
cagcaagggt gctgtgaaa cctggcacca agaccaacat ggtgatggtt ttcggagaga 300
tcacaaccaa ggccaagtgg actatgagaa gat 333

<210> 1993
<211> 325
<212> nucleic acid
<213> Glycine max
<400> 1993

tcgcatgcac gcgtacgtaa gctcgggaatt cggctcgagg ttgagaccaa gacacactcg 60
 ttcatatatc tctctgtctt tctcttctct tctacctctc aagtttttga agtataaaga 120
 tggcagagac attcctattc acctcggagt cagtgaacga gggacaccct gataagctct 180
 gcgaccaatc tccgatgctg tctcgcacgc ttgcctcgaa caggacccag acagcaaggt 240
 tgccctgcga acatgcacca agaccaactt ggtcatggtc ttccggagaga tcaccaccaa 300
 ggccaacgtt gactacgaga agatc 325

<210> 1994
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 1994

acgtcgcatt cagcgcgtacg taagctcggg attcggctcg aggttactgt ctgttcaagc 60
 taccatctct ctctctcttt cttagtgcct ccttgccaga agtaaaaatg gcccaagaaa 120
 ctttctctatt cacatctgaa tcagtgaacg aggggacccc tgacaagctc tgtgaccaga 180
 tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag gttgcctgtg 240
 aaacctggca ccaagaccaa catggtgatg gttttcggng agatcacaac caaggccaag 300

<210> 1995
 <211> 322
 <212> nucleic acid
 <213> Glycine max

<400> 1995

gcgtacgtaa gctcgggaatt cggctcgagn aagcnccact tcaaccacca cacnactctc 60
 tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga cattcctatt 120
 tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgacaaa tctccgatgc 180
 tgtcctcgac gottgccttg aacaggaccc agacagcaag gttgcctgcg aaacatgcac 240
 caagaccaat tgggtcatggt cttcggagag atcaccacca aggccaagtt gactacgaga 300
 agatcgtgcg tgacactgca gg 322

<210> 1996
 <211> 321
 <212> nucleic acid

<213> Glycine max

<400> 1996

tgggagtttag gttctgcacg ctctgcttcc agcgagtgtt ctttcttctgt ttcaacacct 60
taatttgcac acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca 120
tctgaatctg taaacgaggg tcaccccgac aagcgtgcga ccagatctct gatgcagtgc 180
tcgatgcgtg ccttgaacag gacctgaca gcaagggtgc ctgtgagaca tgcaccaaga 240
ccaacatggg catgggtctt ggagagatca caaccaaggc aacgtagata tgagaagatg 300
tcgtgnacat gcgcgaattg g 321

<210> 1997

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 1997

tcgcangcac gcgtacgtaa gctcgggaat tcggctcgan ctgcagccga ttcggctcga 60
ggccttttca gggaaggacc ctaccacngg ttgacagaag tgaccgccta tattgtaagg 120
cagctgcaa agagtgttcg tgggcaaatt gccttgntag aagggtgcatt gtgcaagttt 180
cctatgccat tgggtgtccct gagcccttgt cagtgtttgt ggacncttat ggaactggga 240
agattcctga caaggagatt ctgcaaattg tgaaggagaa ttctgacttc agacctggna 300
tga 303

<210> 1998

<211> 328

<212> nucleic acid

<213> Glycine max

<400> 1998

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ncagacttaa caacagcaca 60
aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180
acctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
acctgacag caagggtgcc tgtgaaacct gcaccaagac caacatggtg atgggtttctg 300

gagagatcac naccaaggcc aacgtggg

328

<210> 1999

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 1999

tcgcanaent acgcncggaa tctcggcncg anaacagcac aaagcgggtt actgtctgtt 60

caagctacca tctctctctc tctntcttag tgctccctt gccagaagt aaaatggccc 120

aagaaacttt cctattcaca tctgaatcag tgaacgagg gcaccctgac aagctctntg 180

accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaangttg 240

cctgtgaaac ctgcaccaag accaactgg tgatngtttt cggagagatc acaaccaagg 300

cnccg 305

<210> 2000

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2000

gtgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60

ggttactgtc tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa 120

gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180

gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240

gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgt tttcggagag 300

atcacaacca ggccaagtgg a 321

<210> 2001

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2001

gtcgcatgca cgcgtacgta agctcgggaa ttcggctcga ggtgatttgg gagtttggag 60

cgactgaact aatcattaat ttgcactcgc tgtttcagct tcatcaccct tcttttgcac 120

catttatatc tcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180
cgaggggtcac cccgacangc tgttcnanca gatctctgat gcagtacttg atgcgtgcct 240
tgaacaggac cctgacagca aggttgacctg tgagacatgc accaagacca acatgggtcat 300
ggtcttcgga gagatcacia ccaaggc 327

<210> 2002
<211> 316
<212> nucleic acid
<213> Glycine max
<400> 2002

ntcgnatnnc agentangtn agnnttcggn tgcgatttg ggagttaggt tctgcacgct 60
ctgcttccag cgagtgttct ttcttcgttt caacacctta atttgcacac gctgctttct 120
tengcttgag aaatgggaca agnnaccttt ctattcacat ctgaatctgt aaacgaggggt 180
caccocgaca anctgtgcga ccagatctct gatgcagtgc tcgatgcgtg ccttgaacag 240
gacnctgaca gcaaggttgc ctgtgagaca tgcaccaaga ccaacatgggt catgggtcttt 300
ggnagatca caacca 316

<210> 2003
<211> 334
<212> nucleic acid
<213> Glycine max
<400> 2003

ccnncngccn acccntngnc nncntntcng tcgnngnnnc gtacgtnagc tcggnattcg 60
gctcngcca agccccactc aaccaccaca ccactctctc tgctcttctt ctacctttca 120
ngtttttaaa gtattaagat ggcagagaca ttctattta cctcagagtc agtgaacgag 180
ggacaccctg acaagctctg cgaccaaata tccgatgctg tctcgcagc ttgccttgaa 240
caggacccag acagcaaggt tgctgcgaa acatgcacca agaccaactt ggcatgggtc 300
ttcggagaga tcaccaccaa ggccaacgtt gact 334

<210> 2004
<211> 216
<212> nucleic acid
<213> Glycine max

<400> 2004

tagtgectcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttggag caggaccctg acagcaaggt tgctgtgaa acctggcacc aagaccaaca 180
tggtgatggg tttcgggagag atcacaacca aggcca 216

<210> 2005

<211> 319

<212> nucleic acid

<213> Glycine max

<400> 2005

gnacgacgca ngcacgcgta cgtnagctcg gaattcggct cgagctcatg gtgatgctgg 60
tctcactgga agaaagatca tcattgatac ctatggtggg tggggtgctc atggtggagg 120
tgcccttttca gggaaggacc ctaccaaggt tgnccagaagt ggtgcctata tcgtgaggca 180
ggctgcaaag agtggttngtg gcaaattggcc ttgccagaag gtgcattgtc caagtttcct 240
atgccattgg gtgtccctga gccctngtca gnggtnggtg gacattatgg nncntgggaa 300
nttccctcaca aggggtttt 319

<210> 2006

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 2006

tcgcatgcac gcgtacgtaa gctcgggaatt cggctcgagc tcgagccgct cgagccgatt 60
cggctcgagg tggccctcat ggtgntgctg gtctcactgg accgaaagat acntcattga 120
tacctatggg ggggtggggac ctcatgggtg aggtgccttt tcaggggaagg accctaccaa 180
ggttgacaga agtggtgcct atatngtgag gcaggctgca aanagtgttg tggcaaattg 240
ccttgccaga aggtgcattg tccaagtttc ctatgcnatt ggtgtccctg agccc 295

<210> 2007

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2007

tatctctctg ttctcttcta cctctcaagt ttttgaagta tagagatggc agagacattc 60
ctatttacct cagagtcagt gaacgaggga caccctgaca agctctgtga ccaaattctct 120
gatgctgtcc tcgacgcttg cctcgaacag gaccagaca gcaagggttg ctgcgaaaca 180
tgcacaaaaa ccaacttggc catggtcttc ggagaatcac gaccaaggcc aatgtngant 240
acgagaagat atgcgtgacc c 261

<210> 2008

<211> 422

<212> nucleic acid

<213> Glycine max

<400> 2008

caggcaagcn ccaactcaacc accacacctc tctngttca cgtacccgc tttctgctct 60
ttttctacct ttcaagtttt aaaagtataa agatggcaga gacattccta ttacctcag 120
agtcggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctng 180
acgcttgctt cgagcaggac ccanacagca aagttgcctg cgaaacatgc accaaaacca 240
acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gaagaagata 300
gtgcgtgaca cctgcaggaa ccattngnnt tngtctnaaa tgatgtgggg actggatgcc 360
cgacaactgg caaggtcctc gtcnaacatt gancatcaaa agccctggtn ttggttnagg 420
gg 422

<210> 2009

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 2009

tcgngcacn cgtacgttag ctcggnnttc ggctcgacct cgagccgaat cggctcgagg 60
ggttactgtc tgttcaagct aaccatctct ctctctctac tontagtgcc tcttgccan 120
aagttaaaat ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180
ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgttg gagcaggacc 240
ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300

agatcacaa

309

<210> 2010
<211> 280
<212> nucleic acid
<213> Glycine max

<400> 2010

ttcaangcag cgtacgtaag ctcggaattc ggctcgagcg gctcgagctc ttctacctct 60
caagtttttg aagtataaag ntggcagaga cattcctatt cacctcggag ttagtgaacg 120
agggacaccc tgataagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctcg 180
aacaggaccc agacagcaag gttgcctgcg aaacatgcac caagaccanc ttggatcatgg 240
nnttcggaga gatcaccacc aaggccaacg ttgactacga 280

<210> 2011
<211> 313
<212> nucleic acid
<213> Glycine max

<400> 2011

atgcacgcgt acgtaagctc ggaattcggc tcgaggcaga cttacaaca gcacaaagcg 60
ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgctc cttgccagaa 120
gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttga gcaggaccct 240
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg ttcgggngng 300
atcanaacaa ggg 313

<210> 2012
<211> 290
<212> nucleic acid
<213> Glycine max

<400> 2012

gtcgcangca cgcgtacgtn agctcggat tcggctcgag gcgggttact gtctgttcaa 60
gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa tggccaaga 120
aactttccta ttcacatctg aatcagtga cgaggggcac cctgacaagc tctgtgacca 180

gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca aggttgcttg 240
 tgaaacctgc accaagacca acatggtgat ggttttcgga gagatcacia 290

<210> 2013
 <211> 274
 <212> nucleic acid
 <213> Glycine max
 <400> 2013

agtcgcannc acgcgtacgt aagctcggaa ttcggctcga nggctcgagc ggctcgnngc 60
 acggccacct cncaaagagg cctgaggaga ttggtgctgg tnnccaaggt catatgttcg 120
 gctatgccac tgacgagact nccgagctca tnncttgag cnatgtcctt gccacnaagc 180
 tccgtgccaa gctcaccgag gttcggaaga acgggacatg cccttggttg agacctgatc 240
 gcaagacca ctccactgtt gagtactaca acgn 274

<210> 2014
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 2014

gtcgcangca cgcgtacgta agctcggaa ttcggctcgag gcacaaagcg ggttactgtc 60
 tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa gttaaaatgg 120
 cccaagaaac ttctctatct acatctgaat cagtgaacga ggggcaccct gacaagctct 180
 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240
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<210> 2015
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 2015

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 atggggccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180

gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240
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 aaccaaggc 309

<210> 2016
 <211> 305
 <212> nucleic acid
 <213> Glycine max
 <400> 2016

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 aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctgaca 180
 agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240
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<210> 2017
 <211> 294
 <212> nucleic acid
 <213> Glycine max
 <400> 2017

gnngcaggcg tacgtaagct cggaattcgg ctcgaggcac aaagcggggt actgtctgtt 60
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 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caaggttgcc 240
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<210> 2018
 <211> 321
 <212> nucleic acid
 <213> Glycine max
 <400> 2018

tcncangcac gcgtacgtaa gctcgggaat tcgggctcga gggcagactt aacaacagca 60

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gccgcaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg 180
gcaccctgac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagca 240
ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaacaagg tgatgggttt 300
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<210> 2019
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<211> 310
<212> nucleic acid
<213> Glycine max
<400> 2020

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taccttgatg agaagacat tttccacttg aaccctctg gccgttttgt cattggaggt 180
cctcacggtg atgctggtct caccggccgc aagatcatca tcgatactta cggaggatgg 240
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ttacatgtga 310

<210> 2021
<211> 326
<212> nucleic acid
<213> Glycine max

<400> 2021

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tccnattttac ctcagagtcg gtgaacgagg gacaccctga caagctctgc gaccaaattct 180
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catgcaccaa aaccaacttg gncatggtct tcggaganat cagaccaag gccaacgttg 300
actacgagaa atagtgcgtg acacct 326

<210> 2022

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2022

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gaatggcacc tgtgtgtggtt gaggccagat ggtaagacac aagtaaccgt cgagtactac 120
aatgacaatg gtgccatggt tccagttcgt gtccacactg tctaatttc cacccaacat 180
gatgagnctg tgagcaatga tcaaagtctg cggaccttaa agagcatggt atcaagcctg 240
tcatactgag aagtaacctg atgagaagac catcttcac ctaaacctc tggccgttt 299

<210> 2023

<211> 545

<212> nucleic acid

<213> Glycine max

<400> 2023

aactgtcnnn cngcgacgcc ngtagctacc agtcactaa tnccggggtc cacncacgn 60
tccgtacggc tgcgaagaag acgacagaag ggganactta ctagtnntaa gcatggagtc 120
ctcaantntc tntgcttntn tcttnanctt naagtttttt aaagtattaa gatggcaaga 180
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aacgttgact angagaaaga tngtgcgtga cacctgcagg aatatcggct tcgtctcagg 420

angatntggg acttgatnct gacatctgca angtecttgt aaacattncg cagcatancc 480
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 ggagg 545

<210> 2024
 <211> 271
 <212> nucleic acid
 <213> Glycine max
 <400> 2024

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 aaagtattaa gatggcagag acatttctat ttacctcaga gtcagtgaac gagggacacc 120
 ctgacaagct ctgcgaccaa actccgatgc tgtctctgac gcttgcccttg aacaggaccc 180
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 gatcaccacc aaggccaacg ttgactacga g 271

<210> 2025
 <211> 297
 <212> nucleic acid
 <213> Glycine max
 <400> 2025

gcaagcccca ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt 60
 ctaacttttc aagtttttaa agtataaaga tggcagagac atttctatctt acctcagagt 120
 cgggtgaacga gggacaccct gacaagctct gcgaccaa atccgatgct gtcctcgacg 180
 cttgcctoga gcaggaccca gacagcaaag ttgcctgcga aacatgnacc aaaaccaact 240
 tgggtcatggt cttcggagaa atcacgacca aggccaagtt gactacgaga agatagt 297

<210> 2026
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2026

gtgcgacgca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60
 gggttactgtc tgttcaagct accatctctc tctctctttc ttagtgctc cttgccagaa 120

gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacaacca 310

<210> 2027
<211> 310
<212> nucleic acid
<213> Glycine max
<400> 2027

gtcgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60
ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgctc cttgccagaa 120
gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacaacca 310

<210> 2028
<211> 309
<212> nucleic acid
<213> Glycine max
<400> 2028

nngnancctta gagtcgcatg cagcgtacg taagctcgga attcggctcg aggttagggt 60
ctgcacgctc tgcttcacgc gagtggtctt tcttcgtttc aacaccttaa tttgcanacg 120
ctgcttcttc ngcttgagaa atggcacaag aaacctttct attcacatct gaatctgtaa 180
acgaggggtca ccccgacaag ctgtgcgacc agatctctga tgcagtgtc gatgcgtgcc 240
ttgaacagga ccttgacagc aaggttgctt gtgagacatg caccaagacc aacatgggtca 300
tggtctttg 309

<210> 2029
<211> 487
<212> nucleic acid
<213> Glycine max

<400> 2029

aactctactt ggncaggccc cggtnacagaa atccccggctc gacccacgcg tcagtacggc 60
tgcgagaaga cgacagaagg gggcagcgct tgatttgagg ccaggcaagc cccactcaac 120
caccacacct ctctcgttc acgtacccc tttctgtctt tttctacct ttcaagtttt 180
aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa cgagggacac 240
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcangac 300
ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 360
gaaatcacga acaagggcaa cgttgactac gaaaaagata attgcntgac aacctgcagg 420
gaacatcggc ttcgtctcaa atgatgttgg gactggatgc cgacaactgc aaaggtctcc 480
gtcaaca 487

<210> 2030

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2030

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gccaaagccc actcaaccac 60
cacaccactc tctctgtctt tttctacct ttcaagtttt taaagtatta agatggcaga 120
gacattccta tttacctcag agtcagttaa cgagggacac cctgacaagc tctgcgacca 180
aatctccgat gctgtcctcg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240
cgaaacatgc accaagacca acttggtcat ggtcttcgga gagatcacca ccaaggcc 298

<210> 2031

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2031

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cagcttgaga aatggcacia gaaacctttc tattcacatc tgaatctgta aacgagggtc 180
accccgacaa gctgtgcgac cagatctctg atgcagtgtc cgatgcgtgc cttgaacagg 240

accctgacag caaggttgcc tgtgagacat gcaccaagac caacatggtc atgggtctttg 300
g 301

<210> 2032
<211> 297
<212> nucleic acid
<213> Glycine max

<400> 2032

cgtcgcangc acgcgtacgt nagctcggna ttccggctcga gngcacaaag cgggttactg 60
tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat 120
ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagct 180
ctgtgaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa 240
ggttgctgt gaaacctgca ccaagaccaa catggtgatg gttttcggng agatcac 297

<210> 2033
<211> 332
<212> nucleic acid
<213> Glycine max

<400> 2033

tgcacgcgta cgtaagctcg gaattcggct cgagatttga ggcaggcaa gcccnactca 60
accaccacac ntctctctgt tnanctacc ctttctncc tcttcttcta cttttcaagt 120
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 180
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgctng cctcgagcag 240
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ggaganatca cgaccaaggc caacgttgac ta 332

<210> 2034
<211> 300
<212> nucleic acid
<213> Glycine max

<400> 2034

tcgcatgcac gcgtacgtna gctcggatt cggtcagagn acagcacaaa gcgggttact 60
gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120

tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180
 tctgtgacca gatctccgat gctgtgntcg atgcatgctt ggagcaggac cctgacagca 240
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<210> 2035
 <211> 307
 <212> nucleic acid
 <213> Glycine max
 <400> 2035

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 gttaaaatgg cccaagaaac tttcctatcc acatctgaat cagtgaacga ggggcaccct 180
 gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
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 atcacia 307

<210> 2036
 <211> 262
 <212> nucleic acid
 <213> Glycine max
 <400> 2036

ccaagcccca ctcaaccacc acaccactct ctctgctctt cttctacctt tcaagttttt 60
 aaagtattaa gatggcagag acattcctat ttacctcaga gtcagtgaac gagggacacc 120
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctt gaacaggacc 180
 cagacagcaa ggttgcttg gaaacatgca ccaagaccaa cttggctcatg gtcttcggag 240
 agatcaccac caaggccaac gt 262

<210> 2037
 <211> 323
 <212> nucleic acid
 <213> Glycine max
 <400> 2037

aaatntanan gtcgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca 60

gcacaaagcg ggttactgtc tgttcaagct accatctcct ctctctcttt cttagtgcct 120
ccttgccaga agttaaagt gccaagaaa ctttcctatt cacatctgaa tcagtgaacg 180
aggggcaccc tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg 240
agcaggaccc tgacagcaag gttgcctgtg aaacctgcac caagaccaac atggtgatgg 300
ttttcggaga gatcacaacc aag 323

<210> 2038
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 2038

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60
aagcgggtta ctgtctgttc aagctaccat ctctctctct cttctcttagt gcctccttgc 120
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttgc 300
gagagatcac a 311

<210> 2039
<211> 301
<212> nucleic acid
<213> Glycine max

<400> 2039

ttcangcacn cgtacgtaag ctcggaattc ggctcgagca caaagcgggt tactgtctgt 60
tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt aaaatggccc 120
aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac aagctctgtg 180
accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaaggntg 240
cctgtgaaac ctgcaccaag accaacatgg tgatggtttt cgngagatc acaaccaagg 300
n 301

<210> 2040
<211> 307
<212> nucleic acid

<213> Glycine max

<400> 2040

gtngcangca cgcgtacgta agctcggaat tcggctcgag cagcaciaag cgggttactg 60
tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat 120
ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagcc 180
ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240
ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaac 300
caaggcc 307

<210> 2041

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2041

cgcatgcagt ntacgtaagc tcggaattcn gctcgagcag caciaagcgg gttactgtct 60
gttcaagcta ccattctctc ctctctttct tagtgctcc ttgccagaag ttaaaatggc 120
tcaagaaact ttctatttca catctgaatc agtgaacgag gaccacctg acaagctctg 180
tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggacctg acagcaaggt 240
tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttcggagaga tcacaacca 300
ggc 303

<210> 2042

<211> 486

<212> nucleic acid

<213> Glycine max

<400> 2042

tngcnaactc ttacgcggtt caggtaccgg ttgnngaatt cccggggtcg acccacgcgt 60
caagtacggc tgcgagaaga cgacagaagg gggcagcgt tgatttgagg ccaggcaagc 120
ccactcaac caccacacct ctctcgttc acgtacccc tttctgctct tttctacct 180
ttcaagtttt aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa 240
cgaggggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgct 300

cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat 360
 ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gaagaagata gtgcgtnaca 420
 cctgcagga acatccgnt nntnccaaaa tnangttgga ncgggatccn anaatttgc 480
 aggggt 486

<210> 2043
 <211> 304
 <212> nucleic acid
 <213> Glycine max
 <400> 2043

ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggggcaccct gnacaagctc 60
 tgtgaccaga tctccgatgc tgtgctcgat ggcattgcttg gagcaggacc ctgacagcaa 120
 ggttgctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaac 180
 caaggccaac gtggactatg agcaagcttg tgnctgaca catgcaggaa cattggtttt 240
 gtctctnatg atgtnggtct tggatgcnaa caactgcaag tctcgtcaac atngagcaac 300
 agan 304

<210> 2044
 <211> 325
 <212> nucleic acid
 <213> Glycine max
 <400> 2044

gtcgcangca cgcgtacgta agctcgaatt cggctcgagg cagacttaac aacagcacia 60
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
 ccctgacagc aaggttgctt gtgaacctgc accaagacca acatggtgat ggttttcgga 300
 gagatcacia ccaggccang tggan 325

<210> 2045
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 2045

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag tgagaaatgg cacaagaaac 60
ctttctattc acatctgaat ctgtaaacga gggtcacccc gacaagctgt gcgaccagat 120
ctctgatgca gtgctcgatg cgtgccttga acaggaccct gacagcaagg ttgcctgtga 180
gacatggcac caagaccaac atggatcatg ttctttggag agatncacaa ccaagggcca 240
acgtagacta tgagaagatt gttcctgnac acatgccggc gaantggatt ncannccg 298

<210> 2046

<211> 318

<212> nucleic acid

<213> Glycine max

<400> 2046

gtcgcacatgca cncgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60
aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
accctgacag caagggttgc tgtgaaacct gcaccaagac caacatggtg aggttttcgg 300
agagatcaca accaaggc 318

<210> 2047

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2047

gngtcgnang cagcgtacg tnagctcgga atgcggctcg aggggttact gtctgttcaa 60
gctaccatct ctncctctct ttcttagtgc ctcttgcca gaagnnaaan tngcccaaga 120
aactttccta ttcnnatctg aatcagtga cgaggggcac cctgacaagc tctgtgacca 180
gatctccgat gctgtgctcg atgcatgcnt ggngcaggac nctgacagca aggttnoctg 240
tgaaacntgc accaagacca acatggtgat ggttttcgga gagatcacia ccaaggccaa 300
cg 302

<210> 2048

<211> 301
 <212> nucleic acid
 <213> Glycine max

 <400> 2048

 tcgnangcac gcgtacgtaa gctcggaatt cggctcnagt ttgggagtta ggttctgcac 60
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120
 ottongcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180
 gtcaccccgca caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240
 aggaccctga cagcaagggt gctgtgaga catgcaccaa gaccaacatg gtcatgggtct 300
 t 301

<210> 2049
 <211> 273
 <212> nucleic acid
 <213> Glycine max

 <400> 2049

 tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc tctctgctct tctcttctct 60
 tctacctctc aagtttttga agtataaaga tggcagagac attcctattc acctcgaggt 120
 cagtgaacga gggacaccct gataagctct gcgaccaa atccgatgct gtcctcgacg 180
 cttgcctcga acaggaccca gacagcaagg ttgcctgcga aacatgcacc aagaccaact 240
 tggatcatggt cttcggagag atcaccacca agg 273

<210> 2050
 <211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 2050

 tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc tgcacgctct gcttccagcg 60
 agtgttcttt cttcgtttca acaccttaat ttgcanacgc tgctttctct ggcttgagaa 120
 atggcacaag aaacctttct attcacatct gaatctgtaa acgaggggtca ccccgacaag 180
 ctgtgcgacc agatctctga tgcagtgtc gatgcgtgcc ttgaacagga ccctgacagc 240
 aaggttgcct gtgagacatg caccaagacc aacatgggtca tggctcttga gagatcacao 300

ccagggccaa cgt

313

<210> 2051
<211> 312
<212> nucleic acid
<213> Glycine max

<400> 2051

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcacaaag 60

cggttactg tctgttcaag ctaccatctc tctctctett tcttagtgcc tccttgccag 120

aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180

ctgacaanct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240

ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300

agatcacaa ca 312

<210> 2052
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 2052

gcgtacgtaa gctcggaatt cngctcgagg cccactcaa ccaccacacc tctcctcggt 60

cacgtaccc ctttctgctc ttcttctacc tttcaagttt taaaagtata aagatggcag 120

agacattcct atttacctca gagtcggtga acgagggaca ccctgacaag ctctgcgacc 180

aaatctccga tgctgtcctc gacgcttgnc tcgagcagga ccagacagc aaagttgcct 240

gcgaaacatg caccaaaacc aacttggtca tggctctcgg agaaatcacg accaaggcca 300

acgttgat 308

<210> 2053
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 2053

gtcgcngcac gcgtacgtaa gctcggaatt cggtcgagg ttaggttctg cacgtctctg 60

ttccagcgag tggtctttct tcgtttcaac accttaattt gcacacgctg cttcttcagc 120

ttgagaaatg gcacaagaaa cctttctatt cacatctgaa tctgtaaacg agggtcaccc 180
cgacaagctg tgcgaccaga tctctgatgc agtgctcgat gcgtgccttg aacaggaccc 240
tgacagcaag gttgcctgtg agacatgnac caagaccaac atggatcatgg tctttggn 298

<210> 2054
<211> 304
<212> nucleic acid
<213> Glycine max

<400> 2054

nanangangt cgcangcacg cgtacgtgag ctccgnattc ggctcgaggn aagccccact 60
caaccaccac accactctct ctgctcttct tctacctttc aagtttttaa agtattaaga 120
tggcagagac attcctatct acctcagagt cagtgaacga gggacaccct gacaagctct 180
gcgaccaaat ctccgatgct gtcctcgacg cttgccttga acaggaccca gacagcaagg 240
ttgcctgcga aacatgcacc aagaccaact tggatcatgt cttcggagag atcaccacca 300
nggc 304

<210> 2055
<211> 481
<212> nucleic acid
<213> Glycine max

<400> 2055

aaactccacc gccaggtac cggatcaaga attcccggtt cgaccacgc gtcnggcgag 60
aagacnacag aagggtacgg ctgcgagaag acgacagaag ggtacggctg cgagaanacg 120
acagaaggggt acggctgcga agaagacgac agaaggtac ggctgcgaga agacgacaga 180
agggtacggc tgcgagaaga cgacagaang gtacggctgc gagaagacga cagaaggggg 240
acacttatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 300
tcgacttcag acctggaatg atcaccatta acttgacct taagaggggt ggccataggt 360
tcctcaagac agctgcttat ggacactttg gaagggatga ccctgacttc acctgggaag 420
ttgtgaagcc actcaantct gaaaaacctc caacctaga atggttgtna atttaancnc 480
c 481

<210> 2056

<211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 2056

 nacgtcgcat gcacgcgtac gtaagctcgg aattcggctc gagtaacaac agcacaaagc 60
 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ctttgccaga 120
 agttaaaatg gcccaagaaa ctttctctatt cacatctgaa tcagtgaacg aggggcaccc 180
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttcg agcaggaccc 240
 tgacagcaag gttgcctnt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300
 agatcacaac caa 313

<210> 2057
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 2057

 tcncatgcac gcgtacgtaa gctcggaatt cggtcgcagg ttactgtctg ttcaagctac 60
 catctctctc tctctttctt agtgctctct tgccagaagt taaaatggcc caagaaactt 120
 tcctattcac atctgaatca gtgaacgagg ggcaccctga caagctctgt gaccagatct 180
 ccgatgctgt gctcgatgca tgcttgagac aggaccctga cagcaagggt gcctgtgaaa 240
 cctgacacca agaccaacat ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg 300
 gatatg 306

<210> 2058
 <211> 325
 <212> nucleic acid
 <213> Glycine max

 <400> 2058

 angcacncgt acgtnagctc ggnattcggc tcgagncana cttaacanca gcacaaagcg 60
 ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgcctc cttgccagaa 120
 gttaaaatgg cccaaganac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
 gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttga gcaggaccct 240

gacagcaagg ttgcctgtga aacctggcac caagaccaac atggtgatgg ttttcggaga 300
 gatcacaacc aaggccaagt ggata 325

<210> 2059
 <211> 286
 <212> nucleic acid
 <213> Glycine max
 <400> 2059

tcgcatgcac gcgtacgtna gctcggaatt cggctcgagc tttctgctct tttctacct 60
 ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag antcggtgaa 120
 cgagggacac cctgacaagc tetgcgacca aatctccgat gctgtcctcg acgcttgctt 180
 cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggttca 240
 tgggtcttcgg agaaatcacg accaaggcca acgttgacta cgagaa 286

<210> 2060
 <211> 280
 <212> nucleic acid
 <213> Glycine max
 <400> 2060

gtcgcangca cgcgtacgta aagctcggaa ttcggctcga gnaaagatgg cagagacatt 60
 cctatattacc tcagagtcgg tgaacgaggg acaccctgac caagctctgc gaccaaattct 120
 ccgatgctgt cctcgacgct tgctcgcagc aggncccaga tagcaaagtt ncntgcgana 180
 catgcaccan aaccnncttg gtcattggtct tcggagnnat cagcaccang gcnancgttg 240
 actanganan gatantgngt gacacctnca ggnacatcgg 280

<210> 2061
 <211> 324
 <212> nucleic acid
 <213> Glycine max
 <400> 2061

gtcgcattgca cgcgtacgta agctcggaa ttcggctcgag gtgatttggg gtttgagcgg 60
 actgaactaa tcattaattt gcactcgcgtg tttcagcttc atcacccttc ttttgcattca 120
 tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180

agggtcaccc cgacaagctg tgcnaccaga tctctgatgc agtacttgat gcgtgccttg 240
ancaggaccc tgacagcaag gttgcctgtg agacatgcac cnagaccaac aggtcatggt 300
cttcggagag atcacaacca aggc 324

<210> 2062
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2062

ganacgtaag ttagctcgga attcggctcg agncttaaca acagcacaaa gcgggttact 60
gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120
tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180
tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240
aggttgcttg tgaaacctgc accaagacca acatggtgat ggtttcggag agatcacaac 300

<210> 2063
<211> 227
<212> nucleic acid
<213> Glycine max
<400> 2063

ntcgcanaca cgcgtacgtn agcncggaat tcggctcgag gtggcaaattg gccttnccag 60
aaggtgcatt gtccaagttt cctatgccat tgggtgccct gagcccttgt cagtgtttgt 120
ggacacttat ggaactggga agattcctga caaggagatt cttcaaattg tgaaggagaa 180
tttcgacttc agacctggaa tgatcaccat taacttggn c ttaaann 227

<210> 2064
<211> 313
<212> nucleic acid
<213> Glycine max
<400> 2064

tnntcgaan nnangctnac ntnagaatgn nntcgannc aagcnccant caaccancac 60
acntctcttc gttcacgcta cccctttctg gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacaccctga 180

caagctctgc gaccaaactct ccgatgctgt cctcgacgct tgccctcgagc aggacccaga 240
cagcaaagtt gccctgcgaaa catgcaccaa aaccaacttg gtcatgggtct tcggagaaat 300
cacgaccaag gcc 313

<210> 2065
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 2065

nttgcannca cncgtacgtn agctcggnan tcggctcgag ncagacttaa caacagcaca 60
nagcgggtta ctntctgttc aagctaccat ctctctctct ctttcttagt ggccctccttg 120
ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180
caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240
gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggg gatggttttc 300
ggagagatca n 311

<210> 2066
<211> 317
<212> nucleic acid
<213> Glycine max

<400> 2066

cgcatgcata agtacgtaag ctcggaattc ngctcgagca agccccactc aaccaccaca 60
cctctcctcg ttcacgctac ccctttctgc ttttcttcta cttttcaagt tttaaaagta 120
taaagatggc agagacattc ctatttacct cagagtcggt gaacaagggg caccctgaca 180
agctctgcga ccaaactctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt catggtcttc ggagaaatca 300
cgaccaaggc caacgtt 317

<210> 2067
<211> 306
<212> nucleic acid
<213> Glycine max

<400> 2067

agtcgcatgc acgcgtacgt aagctcggaa ttcggtctga gacttaacaa cagcacaaag 60
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120
 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240
 ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300
 agatca 306

<210> 2068
 <211> 320
 <212> nucleic acid
 <213> Glycine max

<400> 2068

ancagtcgna tgcacgcgta cgtaagctcg gaattcgget cgagccccac tcaaccacca 60
 caactctect cgttcacgct acccctttct gctcttcttc tacctttcaa gtttttaaaan 120
 tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 180
 caagctctgc gaccaaattc ccgatgctgt cctcgacgct tgccctcgagc aggaccacaga 240
 cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatggtct tcggagaaat 300
 cacgaccaag gccaaagtga 320

<210> 2069
 <211> 318
 <212> nucleic acid
 <213> Glycine max

<400> 2069

nngngcatgc acgcgtacgt nagctcggaa ttcggtctga gcaagcccca ctcaaccacc 60
 acacctctcc tcgttcacgc tacccttttc tgctcttctt ctacctttca agtttttaaaa 120
 gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 180
 acaagctctg cgaccaaatt tccgatgctg tcctcgacgc ttgcctcgag caggaccacg 240
 acagcaaagt tgctgcgaaa acatgcacca aaaccaactt ggtcatgggtc ttoggagaaa 300
 tcacgaccaa ggccaagt 318

<210> 2070

<211> 302
 <212> nucleic acid
 <213> Glycine max

 <400> 2070

 ngtcgcangc acgcgtacgt aagctcggaa ttcggctcga gacttaacaa cagcacaaaag 60
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120
 aagttaaaat ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300
 ag 302

<210> 2071
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2071

 gnanatgcac gcgtacgtaa nctcggaatt cggctcgagt tgggagttag gttctgcacg 60
 ctctgcttcc agcgagtgtt ctttcttctg ttcaacacct taatttgcac acgctgcttc 120
 ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg taaacgaggg 180
 tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt gccttgaaca 240
 ggacctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg tcatggtc 298

<210> 2072
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2072

 cgcangcacg cgtacgtaag ctcggaattc ggetcgaggc agacttaaca acagcacaaa 60
 gcgggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctctttccc 120
 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
 ccctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300

agagatcacn

310

<210> 2073
<211> 289
<212> nucleic acid
<213> Glycine max

<400> 2073

agtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggggttactg tctgttcaag 60
ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat gggcccaaga 120
aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc tctgtgacca 180
gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca aggttgccctg 240
tgaaacctgc accaagacca acatgggtgat gggttttcgga gagatcaca 289

<210> 2074
<211> 309
<212> nucleic acid
<213> Glycine max

<400> 2074

tgcangnan gcgtacgtaa gctcggaatt cggctcgagn cagacttaac aacagcacia 60
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
ccctgacagc aaggttgccct gtgaaacctg caccaagacc aacatgggtga tgggttttcgg 300
agagatcac 309

<210> 2075
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 2075

gtcnngcac gcgtacgtaa gctcggaatt cggctcgagc agacttaaca acagcaciaa 60
gcgggttact gtctgttcaa gctaccatct ctctctctct tttcttagtg ctccttgcca 120
gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180

cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgott ggagcaggac 240
cctgacagca aggttgcttg tgaaacctgc accaagacca acatggtgat ggttttcggg 300
gagatcac 308

<210> 2076
<211> 310
<212> nucleic acid
<213> Glycine max
<400> 2076

gtcgcatgca cgcgtacgtn agctcggant tcggtctgag cttacaaca gcacaaagcg 60
ggttactgtc tgttcaagct acctctctc tctctcttct ttngtgctc cttgccagaa 120
gttaaaatgg cccaaganac tttctattc acntctgant cngtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacaacca 310

<210> 2077
<211> 310
<212> nucleic acid
<213> Glycine max
<400> 2077

cnnatgcacg cgtacgtaag ctccgctcga gccgaatcgg ctcgagggttg agaccaagac 60
aactcgttc atatatctct ctgctcttct cttactcttc tacctctcaa gtttttgaag 120
tataaagatg gcagagacat tcctattcac ctccgagtcg gtgaacgagg gacaccctga 180
taagctctgc gaccaaattc ccgatgctgt cctcgacgct tgctcgaac aggaccaga 240
cagcaagggt gcctgcgaaa catgcaccaa gaccaacttg gtcatggtct tcggagagat 300
caccaccaag 310

<210> 2078
<211> 325
<212> nucleic acid
<213> Glycine max
<400> 2078

<213> Glycine max

<400> 2081

nnangcacgc gtacgtaagc tcggaattcg gctcgagcag acttaacaac agcacaaagc 60
gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ctttgccaga 120
agttaaaatg gcccaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180
tgacagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacaacca aggcca 316

<210> 2082

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2082

ncgtcgcatg cagcggtacg taagctcggg atttcggctc gaggggttac tgtctgttca 60
agctaccatc tctctctctc tttcttagtg ctccttgcc agaagttaa atggccaag 120
aaacttttct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aaggttgcct 240
gtgaaacctg cnaccaagac caacatgggt atggttttcg gagagatcac aaccaangcc 300
aac 303

<210> 2083

<211> 333

<212> nucleic acid

<213> Glycine max

<400> 2083

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag aagccccact caaccaccac 60
acctctctc gtacacgcta cccctttctg ctctctctct acctttcaag ttttaaaagt 120
ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgagg acacctgac 180
aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca ggacctgac 240
agcaaagttg cctgcgaaac atgcacaaa accaattggg catggtcttc ggagaaatca 300

cgaccaaggc caagttgact acgagaagat atg

333

<210> 2084
<211> 287
<212> nucleic acid
<213> Glycine max

<400> 2084

gcacgcgtac gtaagctcgg aattcggctc gaggtgcctt ctctgggaag gatcctacca 60
aggttgatag gagtgggtgcc tacattgtga ggcaagctgc aaagagcatt gttgcaaagt 120
gacttgctag gagggcaatt gtgcaagttt cctatgccat tgggtgtgcct gagccctgtc 180
tgtgtttgtt gacaattatg gcactgggaa gatcccgaca aggaaatcct cagcatgtga 240
aggagagttt tgaactcagcc ggcagatctc catcaacctg atctcaa 287

<210> 2085
<211> 281
<212> nucleic acid
<213> Glycine max

<400> 2085

cgtaacgtaag ctccgaattc ggctcagaca gcacaaagcg ggttactgtc tgttcaagct 60
accatctctc tctctctttc ttagtgcctc cttgccagaa gttaaaatgg cccaagaaac 120
tttctatttc acatctgaat cagtgaacga ggggcacctt gacaagctct gtgaccagat 180
ctccgatgct gtgctcgatg catgcttgga gcaggacctt gacagcaagg ttgcctgtga 240
aacctgcacc aagaccaaca tggatgatgg tttcggagag a 281

<210> 2086
<211> 294
<212> nucleic acid
<213> Glycine max

<400> 2086

gcacgcgtac gtaagctcgg gaattcggct cgaggcagac ttaacaacag cacaagcgg 60
gttactgtct gttcaagcta ccattctctc ctctctttct tagtgcctcc ttgccagaag 120
ttaaaatggc ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccttg 180
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccttg 240

acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcg 294

<210> 2087
 <211> 294
 <212> nucleic acid
 <213> Glycine max

<400> 2087

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ttaacaacag cacaagcgg 60

gttactgtct gttcaagcta ccatctctct ctctctttct tagtgcctcc ttgccagaag 120

ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcacctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggacctg 240

acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcg 294

<210> 2088
 <211> 290
 <212> nucleic acid
 <213> Glycine max

<400> 2088

nngtcgancg cagcgtacg taagctcgga attcggctcg agacagcaca aagcgggtta 60

ctgtctgttc aagctaccat ctctctctct cttcttagt gcctccttgc cagaagttaa 120

aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180

gtctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240

caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggttttcg 290

<210> 2089
 <211> 322
 <212> nucleic acid
 <213> Glycine max

<400> 2089

agtcgcangc ancggtacgt nagctcgga ttcggctcga ggcagactta acaacagcac 60

aaagcgggtt actgtctgtt caagctacca tctctcnctc tctttcttag tgctccttg 120

ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gnacgagggg 180

cacctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg ctcggagcag 240

gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaagn ccaacntggt gatgggttttc 300
ggagannnca anccaagggc an 322

<210> 2090
<211> 318
<212> nucleic acid
<213> Glycine max
<400> 2090

tgcangcac gcgtacgtaa gctcggaatt cggctcgagn ggccaggcaa gccccactca 60
accaccacac ctctctctgt tcacgctacc cttttctgct ottttcttac ctttcaagtt 120
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240
accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300
gagaaatcac gaccaagg 318

<210> 2091
<211> 301
<212> nucleic acid
<213> Glycine max
<400> 2091

tgcangcac gcgtacgtna gctcggaatt cggctcgagc ttaacaacag caciaagcgg 60
gttactgtct gttcaagcta ccattctctt ctctctttct tagtgcttcc ttgccagaag 120
ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcaccctg 180
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg 240
acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttcgagagaga 300
t 301

<210> 2092
<211> 289
<212> nucleic acid
<213> Glycine max
<400> 2092

gtcgcatgca cgcgtacgtn agctcggaat tcggctcgag ccaagcccca ctcaaccacc 60

acacnactct ctctgctctt cttctacctt tcaagttttt aaagtattaa gatggcagag 120
acattcctat ttacctcaga gtcagtgaac gagggacacc ctgacaagct ctgcgaccaa 180
atctccgatg ctgtctctga cgcttgctt gaacaggacc cagacagcaa ggttgccctgc 240
gaaacatgca ccaagaccaa cttgggtcatg gtcttcggag agatcacca 289

<210> 2093
<211> 309
<212> nucleic acid
<213> Glycine max
<400> 2093

gtgcgatgca cggttacgta agctcggaat tcnctcgag gcccactca accaccacac 60
ctctctctgt tcacgctacc cttttctgct cttcttctac ctttcaagtt ttaaaagtat 120
aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180
gtcttgagac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240
caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg gagaaatcac 300
gaccaaggc 309

<210> 2094
<211> 336
<212> nucleic acid
<213> Glycine max
<400> 2094

tgcgancac gcgtacgtaa gctcggaatt cggctcgagg ggccaggcaa gcccactca 60
accaccacac ctncgctcgt ttcacgctac ccctttctgc tcttcttcta cctttcaagt 120
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240
gaccagaca gcaaagttgc ctgcgaaaca tgcaccanaa ccaacttggt catggtcttc 300
ggagaaatca cgaccaaggc caagttgact acgaga 336

<210> 2095
<211> 202
<212> nucleic acid
<213> Glycine max

<400> 2095

tagtgccctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
ggtgatgggtt ttgggagaga tc 202

<210> 2096

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 2096

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctcgagccgg aacttaacaa 60
cagcacaaaag cgggttactg tctgttcaag ctaccatctc tctctctctt tottagtgcc 120
tccttggcag aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac 180
gagggggcacc ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctt 240
gagcaggacc ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg 300
gttttccggag agatc 315

<210> 2097

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2097

cnaagagtcg catgcacgcg tacgtaagct cggaattcng ctcganggca agccccactc 60
aaccaccaca cctctctctg ttcacgctac ccctttctgc tcttcttcta cctttcaagt 120
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggg gaacgagggg 180
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240
gaccagaca gcaaagttgc ctgcgaaaca tgcaccanna ccaacttggg catggtcttc 300
ggagaaatca cgaccaaggc ca 322

<210> 2098

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 2098

nnnttcnngc actcgtagcn aagctcggaa ttcggtctga ggaccaagcc cactcaacc 60
accacaccac tctntctggc tcttcttcta ctttcaagt tnttaaagta ttaagatggc 120
ngagacagcc ctatttacn cagagtcagt gaacgangga caccctgaca agctctgcga 180
ccaaatctcc gatgctgtcc tcgacgcttg ccttgaacag gaccagaca gcaaggttgc 240
ctgcgaaaca tgcaccaaga ccaacttggc catggtctnc ggagagatca ccaccaaggc 300
caacgtt 307

<210> 2099

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 2099

tatgcntnca cgcgtacgta agctcgagaa ttcggtctga gaggccaggc aagccccact 60
caaccaccac acctctnctc gttcacgcta ccccttaatg ctcttctnct acctttnaag 120
ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
acaccctgac aagctctgcg accaaatctc cgatgctgtc ctgcacgctt gcctcgagcn 240
cgaccagac agcaaagttg cctgcgaaac atgcaccaan accaacttgg tcatggtctt 300
cggagaaatc acgaccaagg cca 323

<210> 2100

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2100

gcnngcacgc gtacgtaagc tcggaattcg gctcgaggtt aggttctgca cgctctgctt 60
ccagcgagtg ttctttcttc gtttcaacac cttaatttgc acacgctgct tcttcagctt 120
gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag ggtcaccctg 180
acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa caggaccctg 240
acagcaaggt tgctgtgag acatgcacca agaccaacat ggtcatggt 289

<210> 2101
 <211> 290
 <212> nucleic acid
 <213> Glycine max

 <400> 2101

 gtcgcangca cgcgtacgta agctcggaat tcggctcgag ccaagcccca ctcaaccacc 60
 acacnactct ctctgctctt cttctacctt tcaagttttt aaagtattaa gatggcagag 120
 acattcctat ttacctcaga gtcagtgaac gagggacacc ctgacaagct ctgcgaccaa 180
 atctccgatg ctgtcctcga cgcttgccctt gaacaggacc cagacagcaa ggttgcctgc 240
 gaaacatgca ccaagaccaa cttgggtcatg gtcttcggag agatcaccac 290

<210> 2102
 <211> 301
 <212> nucleic acid
 <213> Glycine max

 <400> 2102

 ggtngtaagg tcggaattcg gctcgagnac cgatgaaacc cccgagtaca tgcccctcag 60
 ccatgtcctt gcaaccaaac ttggtgctcg cntcacagag gttaggaaga atggnacctg 120
 tgcttggttg aggccagatg gtaagaccaa gtaaccgtng agtactacaa tgacaatggt 180
 gccatggttc cagttcgtgt ccacactgtn ctaatttcca cacaacanaa aanncttana 240
 aannaatgat catattgctg cggacttaaa gagcagttat tnaagcctgt gnatctgaga 300
 a 301

<210> 2103
 <211> 311
 <212> nucleic acid
 <213> Glycine max

 <400> 2103

 acgtcgcatg cgcgcgtacg taagctcgga attcggctcg agcaagcccc actcaaccac 60
 cacaccactc tctctgctct tcttctacct ttcaagtttt taaagtatta agntggcaga 120
 gacattccta ttacctcag agtcagtga cgaggacac cctgacaagc tctgcgacaa 180
 atctccgatg ctgtcctcga cgcttgccctt gaacangacc cagacagcaa ggttgcctgc 240

gaaacatgca ccaagaccaa cttgggtcatg gtcttcggag agatcaccac caaggccaag 300
 ttgactagag a 311

<210> 2104
 <211> 313
 <212> nucleic acid
 <213> Glycine max
 <400> 2104

gnacangcac gcgtacgtaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
 agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180
 cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac 240
 cctgacagca aggttgcttg tgaaacctgc accaagacca acatgggtgat ggttttcggg 300
 gagatcacia cca 313

<210> 2105
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2105

ttncngcacg cgtacgtaag ctcggaattc ggctcgagnt aacaacagca caaagcgggt 60
 tactgtctgt tcaagctacc atctctctct cttttctta gtgctcctt gccagaagtt 120
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180
 aagctctgtg accagatctc cgatgctgtg ctgatgcat gcttggagca ggaccctgac 240
 agcaaggttg cctgtgaaac ctggcaccaa gaccaacatg gtgatggttt tcggagagat 300
 cacaac 306

<210> 2106
 <211> 325
 <212> nucleic acid
 <213> Glycine max
 <400> 2106

agttcanaca gcgtacgana gctcggaant cggctcgagg gccaggcnag cccnatcaac 60

cancacacnt ctctacnct cacgctacnc cttgctgcnc ttncgcgac ntngcaagt 120
nctnaaaagt ataaagatgg cagagacatn cctantnacc ncagagtcgg tgaacgaggg 180
anaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240
ggacccagac agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatggtctt 300
cggagaaatc acgaccaagg ccaac 325

<210> 2107
<211> 294
<212> nucleic acid
<213> Glycine max
<400> 2107

aanngangca cgcgtacgta agctcggaat tcggctcgag caacagcaca aagcnggtta 60
ctgtctgttc aagctaccat ctctctctct tttcttagt gcctccttgc cagaagttaa 120
aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc accctgacaa 180
gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240
caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttcg gaga 294

<210> 2108
<211> 304
<212> nucleic acid
<213> Glycine max
<400> 2108

gtcgcacgca cgcgtacgta agctcggaat tcggctcgag ctcgagccgc aacagcacia 60
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180
cctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
cctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300
agan 304

<210> 2109
<211> 303
<212> nucleic acid
<213> Glycine max

<400> 2109

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gagacttaac aacagcacia 60
agcgggttac tgtctgttca agctacnate tctctctctc tttcttagtg cctccttgcc 120
agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
ccctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300
aga 303

<210> 2110

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2110

ngtngcatgc acgcgtacgt aagctcggaa ttcggctcga gcgagccatt tgggagttag 60
gtttctgacg ctctgcttcc agcgagtgtt ctttcttctg ttcaaacacct taatttgac 120
acgtgcttc ttcagcttga gaantggcac aagaaacctt tctattcaca tctgaatctg 180
taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctgatgcgt 240
gccttgaaca ggaccctgac agcaagggtt cctgtgagac atgcaccaag accaacaagg 300
tca 303

<210> 2111

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2111

acgtcgcatt cacgcgtacg taagctcggaa attcggctcg agatttgga gttaggttct 60
gcacgctctg cttccagcga gtgttctttc ttcgtttcaa caccttaatt tgcatacgt 120
gcttcttcng cttgagaaat ggcaagaagaa acctttctat tcacatctga atctgtaa 180
gagggtcacc ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt 240
gaacaggacc ctgacagcaa ggttgcttgt gagacatgca ccaagaccaa catggtca 298

<210> 2112

<211> 286
 <212> nucleic acid
 <213> Glycine max

 <400> 2112

 nnogcangca cgcgtacgta agctcggaat tcggctcgag agccccactc aaccaccaca 60
 ccaactctctc tgctcttctt ctacctttca agtttttaaa gtattaagat ggcagagaca 120
 ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg cgaccaaadc 180
 tccgatgctg tnnctgacgc ttgccttgaa caggaccag acagcaaggt tgctgagaa 240
 acatgcacca agaccaactt ggtcatggtc ttcgagaga tcacca 286

<210> 2113
 <211> 316
 <212> nucleic acid
 <213> Glycine max

 <400> 2113

 cacgtcgcan gcacgcgtac gtaagctcgg aattcggctc gactgacaag gagattctgc 60
 aaattgtgaa ggagaatttc gacttcagac ctggaatgat caccattaac ttggacctta 120
 agaggggtgg tcataggttc ctcaagacag ctgcttatgg acactttgga agggatgatg 180
 cagacttcac ctgggaagtt gtgaagccac tcaagtcaga gaagcctcaa gcttaagagt 240
 gttgttaagt taatcactcc cttcagtggg tgtcttgctg ggtgtggatg aataatttgc 300
 gtgtttcatg actact 316

<210> 2114
 <211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 2114

 tgcangcacg cgtacgtaag ctcggaattc ggctcgagnt atcaagcctg tcattcctga 60
 gaagtacctt gatgacatgc tctttaaggt ccgcagcaat ttgatcattg ctacagtct 120
 ncatcatggt ggggtggacct taaagagcan nttntcaagc ctgtcattcc tgagaagtac 180
 cttgatgaga agaccatctt ccaccttaac ctttctggcc gttttgtcat tgggtggcct 240
 catggtgang ctgcnctcac tggaagaaa atcatcattg atacctatgg tggctggggt 300

gctcatgg

308

<210> 2115
<211> 284
<212> nucleic acid
<213> Glycine max

<400> 2115

ctngcnctng tacgttagct cggaattcgg ctcgaggnac caagccccac tcaaccacca 60
caonactctc tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga 120
cattcctatt tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgaccaa 180
tctccgatgc tgtctcgac gcttgcttg aacaggaccc agacagcaag gttgcctgcg 240
aaacatgcac caagaccaac ttggtcatgg tcttcggaga gatc 284

<210> 2116
<211> 283
<212> nucleic acid
<213> Glycine max

<400> 2116

cgcangcacg cgtacgtaag ctcggaattc ggctcgagcc aagccccact caaccaccac 60
accactctct ctgctcttct tctacctttc aagtttttaa agtattaaga tggcagagac 120
attcctatct acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaa 180
ctccgatgct gtctctgacg cttgccttga acaggacca gacagcaagg ttgcctgcga 240
aacatgcacc aagaccaact ttggtcatgg cttcggagag atc 283

<210> 2117
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 2117

ngtcgcatgc acgcgtacgt aagctcggaa ttccggtcga ggacttaaca acagcacaaa 60
gogggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca 120
gaancgcaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240

ccctgacagc aaggttgcc t gtaaacctg caccaagacc aacatgggtga tggttttc 298

<210> 2118
<211> 288
<212> nucleic acid
<213> Glycine max

<400> 2118

annnaancaa gcgtacgtaa gctcggaatt cggctcgagg ttaggtttctg cacgctctgc 60
ttccagcgag tgttctttct tcgtttcaac accttaattt gcacacgctg cttcttcagc 120
ttgagaaatg gcacaagaaa ctttctatt cacatctgaa tctgtaaacy agggtcaccc 180
cgacaagctg tgcgaccaga tctctgatgc agtgcctgat gcgtgccttg aacaggaccc 240
tgacagcaag gttgcctgtg agacatgcac caagaccaac atgggtcat 288

<210> 2119
<211> 329
<212> nucleic acid
<213> Glycine max

<400> 2119

tgcangcacg cgtacgtaag ctcggaattc nntcagaggc annacnccan tncaaccacc 60
acacctctcn tcgttcangc tannnaaatn ctgctgttct tctacctgac aagttttgaa 120
agtatanaga tggcaganac attcctattt acctcanagt cggatgaacga gggacaccct 180
gacaagctct gcgaccaaatt ctccngtgcg gtcctcgacg cttgcntcga gcagnaccca 240
gacagcaaag ttgcncgcga nacatggacc aaaaccaact tggatcatgg ntccggagaa 300
atcacgacca aggccaaact tgactacnn 329

<210> 2120
<211> 277
<212> nucleic acid
<213> Glycine max

<400> 2120

ggccaggcaa gcccactca accaccacac ctctcctcgt tcacgctacc cttttctgct 60
cttcttctac ctttcaagtt ttaaaagtat aaagatggca gagacattcc tatttnoctc 120
agagtcggtg aacgaggac accctgacaa gctctgcgac caaatctccg atgctgtcct 180

cgacgcttgc ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacccaaaac 240
 caacttggtc atggtcttcg gagaaatcac gaccaag 277

<210> 2121
 <211> 286
 <212> nucleic acid
 <213> Glycine max
 <400> 2121

cgcangcacg cgtacgtaag ctcggaattc ggctcgagca agccccactc aaccaccaca 60
 cgcgtcnctc tngcgcttct tctaccttct aagtttttaa agtattaaga tggcaganac 120
 attcctatctt acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaaact 180
 ctccgatgct gtctcgcagc cttgccttga acaggaccca gacagcaagg ttgcctgcga 240
 aacatgcacc aagaccaact tggatcatggt cttcggagag atcacc 286

<210> 2122
 <211> 339
 <212> nucleic acid
 <213> Glycine max
 <400> 2122

annctgaanc gtangnaagc ntacgnattc ngctcgagng gcaggcaagc cccactcaan 60
 caccacacct gctcctgcgt ncangctnac ccgtnnnngan gnnatgacta cctntcaagt 120
 tntaaaagta tngnanatgg cngagacatt cctatttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaatctc cgntgctgtc ctcgacgctt gcctcgagca 240
 ggaccagac agcaaagttg cctgcgaaac atgcaccacc accaagttgg tcatggtctt 300
 cggagaaatc acgaccaagg cnaacgttac tacgagann 339

<210> 2123
 <211> 480
 <212> nucleic acid
 <213> Glycine max
 <400> 2123

anctcttacc ggctntntng cccaaaatng tanangcttc ccggctcgac ncacgcgtcn 60
 gtacggctgc gagaagacga cagaaggggg cagctcttga tttnaggnca ngcaancccc 120

actcaanac cacacctctc ctcggtcacg ctatcccttt ctgctcttct tctacctttc 180
angttttaan agtacncaca tggcaagaca cattcctatt tancnagac tcggtgaann 240
acggacaccc tgacaagctc tgcgaccaa tctccgatnc tgtcctcgac gcttgccctcg 300
ancaggactc agacancana nttgcctgcn aaacatgcac caaaaccaac ttggtcatgg 360
tcttcngaga antcacgacc aaggccaacn ttgactacga aaaganngtg cgttacacct 420
gcgggaaca tcggcttctt tcnaaatgat gttgggactg gatgccgacc actgcatnng 480

<210> 2124
<211> 307
<212> nucleic acid
<213> Glycine max

<400> 2124

anntgcang cagcgtacg tnagctcgga attcggctcg agccttaaca acagcacaaa 60
gggggttaact gtctgttcaa gctaccatct ctctctctac tttcttagtg cctccttgcc 120
agangttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
ccctgacaag ctctgtgacc agatctccga tgcgtgtgctc gatgcatgct tggagcagga 240
nnctgacag caagggttgcc tgtgaaacct gcaccaagac caacatgggtg atggttttcg 300
gagagat 307

<210> 2125
<211> 307
<212> nucleic acid
<213> Glycine max

<400> 2125

tcgnngacgc gtacgtaagc tcggaattcg gctcgaggac ttaacaacag caciaagcgg 60
gttactgtct gttcaagcta ccctctctct ctncntttc ctnagtgcct ccttgccaga 120
agttaaaatg gcccaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180
tgaaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300
atcacia 307

<210> 2126

<211> 309
 <212> nucleic acid
 <213> Glycine max

 <400> 2126

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 ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120
 aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc accctgacaa 180
 gctctgtnac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240
 caaggttgcc tgtgnaaacc tggcaccaag accaacaatgg tgatggtttt cggagagatc 300
 acaaccaag 309

<210> 2127
 <211> 302
 <212> nucleic acid
 <213> Glycine max

 <400> 2127

 aaaaanntnaa naggctacgt aagctcgga ttcggctcga gnagacacac tcgttcatat 60
 atctctctgc tcttctcttc tcttctacct ctcaagtttt tgaagtataa agatggnaga 120
 gacattccta ttcacctcgg agtcagtga cgagggacac cctgataagc tctgcgacca 180
 aatctccgat gctgtcctcg acgcttgctt cgaacaggac ccaganagca aggttgccctg 240
 cgaaacatgc accaagacca attggatcatg gtcttcggag agatcaccac caaggccaac 300
 gt 302

<210> 2128
 <211> 288
 <212> nucleic acid
 <213> Glycine max

 <400> 2128

 gtcgcangca cgcgtacgtn agctcggaat tcggctcgag cacanagcgg gttactgtct 60
 gttcaagcta ccatctctac tctctctttt ttagtgcttc cttgccagaa gttanaatgg 120
 cccaagaaac tttcctattc acgtctgaat cagtgaacga ggggcaccct gacaagctct 180
 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240

ttgcctgtga aacctgcacc aagaccaaca tggatgatgt tttcggag 288

<210> 2129
 <211> 279
 <212> nucleic acid
 <213> Glycine max
 <400> 2129

gtgcgacga cgcgtacgta agctcggaat tcggctcgag cacactcgtt catatatctc 60
 tctgtctctt tcttctcttc tacctctcaa gtttttgaag tataaagatg gcagagacat 120
 tctatttcac ctggagtgca gtgaacgagg gacacctga taagctctgc gaccaaactc 180
 ccgatgtgt cctcgacgtt tgctcgaaac aggaccaga cagcaagggt gcctgcgaaa 240
 catgcaccaa gaccaacttg gtcattggtc tcggagaga 279

<210> 2130
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 2130

cgctcgatgc acgcgtacgt aagctcgga attcggtcg agcgagccat ttgggagtta 60
 gggtctgcac gctctgcttc cagcgagtgt tcttctctcg tttcaacacc ttaatttgca 120
 caagctgctt cttcagcttg agaaatggca caagaaacct ttctattcac atctgaatct 180
 gtaaacgagg gtcacccgca caagctgtgc gaccagatct ctgatgcagt gctcgatgcg 240
 tgcttgaac aggacctga cagcaagggt gcctgtgaga catgcaccaa gaccaacatg 300
 g 301

<210> 2131
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 2131

gtgcgacga cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60
 aagcgggtta ctgtctgttc aagctacat ctctctctct ctttcttagt gcctccttgc 120
 cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc 180

accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggttttt 299

<210> 2132

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2132

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aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120

cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180

accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggttttc 299

<210> 2133

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2133

gtcgnacaca aagcgtagtn aagctcggaa ttcggctcga gcggagattc ttcaaattgt 60

gaaggagaat ttcgacttca gacctggaat gatcaccatt aacttggacc ttaagagggg 120

tggccatagg ttctcaaga cagctgctta tggacacttt ggaagggatg accctgactt 180

cacctgggaa gttgtgaagc cactcaagtc tgagaagcct caagcttaag attgttgtga 240

agttaatcac tcccttcaat ggatgtcttg ctaggtgtgg atgaataatt tgcgtgttcc 300

atgactacta ctacttcac 320

<210> 2134

<211> 313

<212> nucleic acid

<213> Glycine max

<400> 2134

nnaatnntaa ngcacgnta cgtaagctcg gaattcggct cgaggggagc cccactcaac 60

caccacacct ctctctgttc acgctacccc tttctgctct tcttctacct ttcaagtttt 120

aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa cgagggacac 180
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
 ccagacagca aagttgcctg cgaaatntgc accaaaacca acttggtcat ggtcttcgga 300
 gaaatcacga cca 313

<210> 2135
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2135

angnatcgca ngcncgcgta cgtnagctcg gaattcggct cgagggccag gcaagcccca 60
 ctcaaccacc acacctctcc tcgttcnccg tacccttttc tgctcttctt ctacctttca 120
 ngtttttaaaa gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag 180
 ggacaccctg acaagctctg cgaccaaata tccgatgctg tctcgcagcg ttgcctcgag 240
 caggaccagc acagcaaagt tgcttcgcaa acatgcacca aaaccaactt ggtcatggtc 300
 ttccggagaaa tcacga 316

<210> 2136
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 2136

agtcgcangc acgcgtacgt aagctcggaa ttcnctcga ggcaagcccc actcaaccac 60
 cacacctctc ctggttcacg ctaccctttt ctngctcttc ttctaccttt ccaagtttta 120
 aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgcttc gagcaggacc 240
 cagacagcaa agttgcctgc gaaacatgca ccaaaaccaa cttgggtcatg gtcttcggag 300
 aaatcacga 309

<210> 2137
 <211> 280
 <212> nucleic acid
 <213> Glycine max

<400> 2137

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tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt aaaatggccc 120
aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcacccngac aagctctgtg 180
accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaagggtg 240
cctgtgaaac ctgcaccaag accaacaagg tgatggtttt 280

<210> 2138

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2138

gtcgcangca cgcgtacgta agctcggaat tcggctcgag cggctcgaga tttgggagtt 60
aggttctgca cgctctgctt ccagcgagtg ttctttcttc gtttcaacac ctttaatttg 120
acacgtgct tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc 180
tgtaaacgag ggtcaccctg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc 240
gtgccttgaa caggaccctg acagcaagggt tgctgtgag acatgcacca agaccaacat 300
ggt 303

<210> 2139

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2139

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cgctctgctt ccagcgagtg ttctttcttc gtttcaacac ctttaatttg acacgtgct 120
tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180
ggtcaccctg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240
caggaccctg acagcaagggt tgctgtgag acatgcacca agaccaacat ggt 293

<210> 2140

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 2140

antcncngca cgcgtacgta agctcggaat tcggctcgag tgatttgagg ccaggcaagc 60
cccaactcaac caccacacct ctctcgttc acgctacccc tttctgctct tttctacct 120
ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa 180
cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct 240
cgagcaggac ccagacagca aagtngcctg cgaaanatgc accagaacca acttggtcat 300
ggtcttcgga gaaatcacga ccaag 325

<210> 2141

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2141

ntcgcangca cgcgtacgta agctcggaat tcggctcgag ctgacaagga gattcttcaa 60
attgtgaagg agaattncga cttcanacct ggaatgatca ccattaactt ggaccttaag 120
aggggtggcc ataggttcct caagacagct gcttatggac actttggaag ggatgaccct 180
gaattcacct gggaagttgt gaagccactc aagtctgaga agcctcaagc ttaagattgt 240
tgtgaagtta atcaactcct tcaatggatg tcttgctagg tgtggatgaa taatttgc 298

<210> 2142

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2142

cgctnnacgt cgcangcacg cgtacgtaan ctcggaattc ggctcgagnt tgggagttag 60
gttctgcacg ctctgcttcc agcgagtgtt ctttcttcgt ttcaacacct taatttgcac 120
acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180
taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctogatgcgt 240
gccttgaaca ggacctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg 300
t 301

<210> 2143
 <211> 283
 <212> nucleic acid
 <213> Glycine max

 <400> 2143

 gogtacgtaa gctcggaatt cggctcgagc aacagcacia agcgggttac tgtctgttca 60
 agctaccatc tctctctctc ttctttagtg ctccttgcc agaagttaa atggccaag 120
 aaactttcct attcacatct gaatnagtga acgaggggca ccctgacaag ctctgtgacc 180
 agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aaggttgct 240
 gtgaaacctg caccaagacc aacatggtga tggttttcgg aga 283

<210> 2144
 <211> 293
 <212> nucleic acid
 <213> Glycine max

 <400> 2144

 ncgtcgcatg cagcggtacg taagctcgga attcggtcg agcaacagca caaagcgggt 60
 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180
 aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240
 agcaagggtg cctgtgaaac ctgcaccaag accaactgg tgatggtttt cgg 293

<210> 2145
 <211> 294
 <212> nucleic acid
 <213> Glycine max

 <400> 2145

 gtgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60
 ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgcctc cttgccagtt 120
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180
 aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240
 agcaagggtg cctgtgaaac ctgcaccaag accaactgg tgatggtttt cgga 294

<210> 2146
 <211> 291
 <212> nucleic acid
 <213> Glycine max

 <400> 2146

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 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120
 agttaaaatg gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atgggtgatgg t 291

<210> 2147
 <211> 340
 <212> nucleic acid
 <213> Glycine max

 <400> 2147

 acgogtacgt aagctcggaa ttcggctcga gggccaggca agccccactc aaccaccaca 60
 cntctectgc gttcangcta cccctttctn gctcttcttc tacctntcaa gtnttaaaag 120
 tataaagatg gcagagacat tectatttac ctcagagtcg gtgaacgagg gacaccctgn 180
 caagctctgc gaccaaatct ccgatgctgt cctcgacgct tgccctcgagc aggaccacaga 240
 cagcaaagtt gcttggcgaa acatgcacca ntnnnacttg gtcatgggtct tcggagaaat 300
 caagaccaag gccaacgttg actacgagaa gatagtgcgt 340

<210> 2148
 <211> 319
 <212> nucleic acid
 <213> Glycine max

 <400> 2148

 agnancgtca cgcgtacgta agctcgggaat tcggctcgag atttgaggnc aggcaagccc 60
 cactcaacca ccacacctct cctcgttcac gctaccctt tctgctcttc ttctaccttt 120
 caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180
 agggacaccc tgacaagctc tgcgacaaa tctccgatgc tgtcctcgac gcttgcctcg 240

agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg 300
tcttcggaga aatcacgac 319

<210> 2149
<211> 198
<212> nucleic acid
<213> Glycine max
<400> 2149

tagtgctcc ttgccagaag ttaaaatggc ccaaganact ttctattca natctgaatc 60
agtgaacgag gggcaccttg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttgag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
ggatgatggt ttcggaga 198

<210> 2150
<211> 293
<212> nucleic acid
<213> Glycine max
<400> 2150

ngtencange acgcgtacgt nagctcgga ttcggctcga ggcacaaagn gggttactgt 60
ctgttcaage taccatctct ctctctgctt tgcttagtgc ctcttgcca gaagttaaaa 120
tgccccaaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180
tctgtgacca gatctccgat gctgtgctn ngccatgctt ggagcaggac cctgacagca 240
aggttgcntg tgaaacctgc accaagacca acatgggtgat ggttttcgga gag 293

<210> 2151
<211> 295
<212> nucleic acid
<213> Glycine max
<400> 2151

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ccaaggccaa cgttgactac gagaagatag tgctgacac ctgcaggaa atgcggcttc 120
cgtctcaa at gatgtgggac tggatgccga caactgcaag gtcctcgtca acattgagca 180
gcagagccnt gatattgcct cagggtgtac acggnccacc ttacnnnnaa acctgaagaa 240

nttgggtgctg gtgaccaggg tccacatggt tggctatgcc atgatgaaac ccnc 295

<210> 2152
 <211> 219
 <212> nucleic acid
 <213> Glycine max

<400> 2152

tagtgctcc tcgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
 agtgaacgag gggcaccctg acaagtctgt gaccagatct ccgatgctgt gctcgatgca 120
 tgcttgagc aggaccctga cagcaagggt gctgtgaaa cctggcacca agaccaacat 180
 ggtgatggtt ttgggagaga tcacaancaa ggccaacgt 219

<210> 2153
 <211> 218
 <212> nucleic acid
 <213> Glycine max

<400> 2153

tantgctcc ttgtcagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60
 agtgaacgag gggcaccatc gacaagctct gtgaccagat ctccgatgct gtgctcgatg 120
 catgcttggc gcaggaccct gacagcaagg ttgctgtga aacctgcacc aagnaccaac 180
 atggtgatgg ttttcggaga gatcacaacc aaggccaa 218

<210> 2154
 <211> 291
 <212> nucleic acid
 <213> Glycine max

<400> 2154

cangcgtacg taagctcgga attcggctcg agacagcaca aagcgggtta ctgtctgtnc 60
 aagcnaccat ctncctctct ctttcttagt gctccttgc cagaagttaa aatggcccaa 120
 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180
 cagatctccg atggctgtnc tcgatgcatg cttggagcag gacctgaca gcnaggttgc 240
 ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca n 291

<210> 2155

<211> 309
 <212> nucleic acid
 <213> Glycine max

 <400> 2155

 tncngtnnnn ngcacgcgta cgtaagctcg gaattcggct cgagnactta anaanagcac 60
 aaagcggggtt actgtctgtt caagctanca tctctctctc tctttcttag tgcttccttg 120
 ccagaagtta aaatggggccc aagaaacttt cctattcaca tntgaatcag tgaacgaggg 180
 gcaccctgac aagctctgtg accagatctc cgatgctgtg cttcgatgca tgcttgagac 240
 aggaccctga cagcaagggtt gcctgtgaaa cctgcaccaa gaccaacatg gtgatgggtt 300
 tcggagaga 309

<210> 2156
 <211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 2156

 nantgcgatn cagcggtacg taagctcgag aatggccach cagccccac tcaaccacta 60
 naontntect cnttcacgct acccctttct gctctncttn tacntttcaa gttttaaaan 120
 nataaagatg gcagagacat tcctatttan ctcagagtcg gtgaacgagg gacaccctga 180
 caagctctgc gaccaaactc ccgatgctgt cctcgacgct tgctcgagc aggaccaga 240
 cagcaaagtt gcctgcgana catgccacca aaaccaactt ggtcatggtc ttcgagaaa 300
 tcacgaccaa ggc 313

<210> 2157
 <211> 294
 <212> nucleic acid
 <213> Glycine max

 <400> 2157

 agtcgcangc acgcgtacgt aagctcgga ttcggctcga ggttcatata tctctctgct 60
 cttctcttct cttctacctc ncaagttttt gaagtataaa gatggcagag acattnccta 120
 ttcacctcgg agtcagtga cgaggacac cctgataagc tctgcgacca aatctccgat 180
 gctgtcctcg acgcttgctt cgaacaggac ccagacagca aggttgctg cgaaacatgc 240

accaagacca acttggttca tgggtcttcgg agagatcacc accaaggcca acgt 294

<210> 2158
 <211> 285
 <212> nucleic acid
 <213> Glycine max
 <400> 2158

cnogcacgcg tacgtaagct cggaattcgg ctcgagaaag cgggttactg tctgttcaag 60
 ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat ggcccaagaa 120
 actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag 180
 atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa ggttgccctgt 240
 gaaacctggc accaagacca acatggtgat gggtttcggg gagat 285

CGACGNGTAC

<210> 2159
 <211> 300
 <212> nucleic acid
 <213> Glycine max
 <400> 2159

gcacgngtac gtnagctcgg aattcggctc gaggccattt gggagttagg ttctgcacgc 60
 tctgcttcca gcgagtgttc tttcttcggt tcaacacctt aatttgaca cgctgcttct 120
 tcagcttgag aaatggcaca agaaaccttt ctattcacat ctgaatctgt aaacgagggg 180
 caccctgaca agctgtgcga ccagatctct gatgcagtgc tcgatgcgtg ccttgaacag 240
 gacctgaca gcaaggttgn ctgtgagact gcaccaagac caacatgggc atggtctttg 300

<210> 2160
 <211> 258
 <212> nucleic acid
 <213> Glycine max
 <400> 2160

gtcgcangca cgcgtacgta agctcggaat tcggctcggn ctcgagccga atcggtctga 60
 gccaacgat gaaacccccg agtaactgcc ctcagccat nctccttgca accaaacttg 120
 gngctcgnt cacagaggtt aggaagaatg gcacctgtgc ttggttgagg ccagatggta 180
 agacacaagt aaccgtcgag tactacaatg acaatgggtc catggttcca gttcgtgtcc 240

acactgtcct aatttcca

258

<210> 2161
<211> 335
<212> nucleic acid
<213> Glycine max

<400> 2161

gtcgcangcn cgcgtagctn nagctcggaa ttcggctcga gcttgatttg aggccaggca 60
agccccactc aacnaccaca cctctcctcg ttcacgctac ccctttctgc tcttcttcta 120
ccttcaagtt ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg 180
aacgagggac accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc 240
ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacaaaaac caattggtca 300
tgggtcttcgg agaaatcacg accaaggcca acgtt 335

<210> 2162
<211> 287
<212> nucleic acid
<213> Glycine max

<400> 2162

cgcaggaccc cccacncnag ctcggaattc ggctcgagcc aagncctact caaccaccac 60
accactctct ctgctcttct tctacctttc aagtnggtaa agtattaaga tggcagagac 120
attcctatctt acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaaatt 180
ctccgatgct gtctctgacg cttgccttga acaggaccca gacagcaagg ttgcctgcga 240
aacatgcacc aagaccaact tggatcatggt cttggagaga tcaccac 287

<210> 2163
<211> 319
<212> nucleic acid
<213> Glycine max

<400> 2163

cngtangacg tcgcatgcac gcgtacgtaa gctcggaaat tcggctcgag ngacntaaca 60
acagcacaaa gcgggttact gtctgttcaa gntanccatc tntgctctct ctttcttagt 120
gcctccttgc nagaagntan aatggcccaa gnaaactttc ctattcacat ctgaatcagt 180

gancgagggg caccctgaca agctctgtga ncagatctcc gatgctgtgc tcgatgcatg 240
 cttggagcag gaccctnaca gcaaggttgc ctgtgaaacc tgcaccaaga ccaanatggt 300
 gatngttttc ggagagatc 319

<210> 2164
 <211> 327
 <212> nucleic acid
 <213> Glycine max
 <400> 2164

nagnntntgc acgcgtacgt aagctcggaa ttcggctcna gcacaaagcg gggttactggc 60
 tgtncaaagt accattctct ctctctcttt cttagtgcct ccttgccata agttaaaatg 120
 gcccagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tnacagcaag 240
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg tttcggagag atcacgacca 300
 aggncaantg ggtntgagaa gatngtg 327

<210> 2165
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 2165

gtcgcattgca cgcgtacgta agctcggaa ttcggctcgag ggccaggcaa gcccactca 60
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 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctgagcagg 240
 acccagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300
 gagaaatca 309

<210> 2166
 <211> 260
 <212> nucleic acid
 <213> Glycine max
 <400> 2166

aagccccact caaccaccac acctctctc gttcacgcta cccctttctg ctctttctct 60
 acctttcaag ttttaaaagt ataaagatgg cagagacatt cctattttacc tcagagtcgg 120
 tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt 180
 gcctcgagca ggaccagac agcaaagttg cctgcgaaac atgcaccaa accaacttgg 240
 tcatggtctt cggagaaatc 260

<210> 2167
 <211> 266
 <212> nucleic acid
 <213> Glycine max
 <400> 2167

aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt tctgctcttc 60
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 120
 tgggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac 180
 gcttgccctg agcaggaccc agacagcaa gttggctgcg aaacatgcac caaaaccaac 240
 ttggtcatgg tcttcggaga aatcac 266

<210> 2168
 <211> 313
 <212> nucleic acid
 <213> Glycine max
 <400> 2168

agnogntgca cgcgtacgta agctcggaat tcnctcgag gccaggcaag cccactcaa 60
 ccaccacacc ttctccttcg ttcacgctac ccttttctgc ttcttcttct acctttcaag 120
 ttttaaaagt ataaagatgg cagagacatt cctattttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240
 ggaccagac agcaaagttg cctgcgaaac atgcaccana accaacttgg tcatggtctt 300
 cggagaaatc acg 313

<210> 2169
 <211> 290
 <212> nucleic acid
 <213> Glycine max

<400> 2169

angcacgcgt acgtaagctc ggaattcggc tcgaggccat ttgggagtta ggttctgcac 60
gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120
cttcngcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180
gtcacccega caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240
aggacctga cagcaagggtt gcctgtgaga catgcaccaa gaccaacatg 290

<210> 2170

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2170

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctcgagccga attcggctcg 60
nngagaaatc acgaccangg ccaaanttga ctacgagaag anngtgcttg acacctgcag 120
gancatcggc ntcgtcncaa atgatgtggg actggangcc gacaactgca aggtcctcgt 180
caacatngag cagcanagcc ctganattgc tcagggngta cncggccacc ttaccaaaaa 240
acctgaagaa attggtgcng g 261

<210> 2171

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 2171

gtgcattcgt acgtaagctc ggaattcngc tcgaggccag gcaagcccca ctcaaccacc 60
acacctctcc tcgttcacgc tacccttttc tgctcttctt ctacctttca agttttaaaa 120
gtataaagat ggcagagacn ttctatttta cctcagagtc ggtgaacgag ggacaccctg 180
acaagctctg cgaccaaadc tccgatgctg tctcgcacgc ttgcctcgag caggacccag 240
acagcaaagt tgcttgcgaa acatgcacca aaaccanctt ggtcatggtc ttcggagaaa 300
tcacg 305

<210> 2172

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 2172

tcgcangcac gcgtacgtaa gctcggaatt ctntctgagg caagccccac tcaaccacca 60
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 180
caagctctgc gaccaaatct ccgatgctgt cctcgacgct tgcctcgagc aggaccaga 240
cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatggtct tcggagaaat 300
caga 304

<210> 2173

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2173

ngangcacgc gtacgtaagc tcggaattcn gctcgaggca agccccactc aaccaccaca 60
cctctcctcg ttacgctac ccctttctgc tcttcttcta cctttcaagt tttaaaagta 120
taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga caccctgaca 180
agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
gnaagttgc ctgcgaaaca tgcaccanaa ccaacttggc catggtcttc ggagaaatca 300
cganca 306

<210> 2174

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 2174

nnncanangc acgcgtacgt aagctcgga ttcggctcga gcggctcgag accactctct 60
ctgctcttct tctaccttc aagtttttaa agtattaaga tggcagagac attcctatct 120
accttcagag tcagtgaacg agggacaccc tgacaagctc tgcgacaaa tctccgatgc 180
tgtctctgac gcttgcttg ancaggaccc agacagcaag gttgcctgcg aaacatgcac 240
caagaccaac ttggtcatgg tcttcggaga gatcaccacc aag 283

<400> 2175

[illegible]

<400>	2176
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<400>	2177
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aagctctgtg accagatctc cgatgctgtg ctcgntgcat gcttggagca ggaccctgan 240
 agcaaggttg cctgtgaaac ctggcaccan gaccaacatg gtgatggttt tcggaga 297

<210> 2178
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2178

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 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240
 ccctgacagc aagggttgct gtgaaacctg caccaagcca acatgggtgat ggttttcggg 300
 gagatcacia 310

<210> 2179
 <211> 278
 <212> nucleic acid
 <213> Glycine max
 <400> 2179

cggtacgtn agctcggaat tcggctcgag cttacaaca gcacaaagcg gggttactgtc 60
 tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa gttaaaatgg 120
 cccaagaaac ttctctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180
 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240
 ttgcctgtga aacctgcacc aagaccaaca tggatgat 278

<210> 2180
 <211> 281
 <212> nucleic acid
 <213> Glycine max
 <400> 2180

cgcatgcacg cgtacgtaag ctcggaattc ggctcgagca acagcnaaaa gcgggttact 60
 gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctccttgcca gaagttaaaa 120

tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180
tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240
aggttgcttg tgaacactgc accaagacca acatggtgat g 281

<210> 2181
<211> 305
<212> nucleic acid
<213> Glycine max
<400> 2181

gnnnangcac gcgtacgtna gctcggaatt cggctcgagg gccaggcaag ccccaactcaa 60
ccaccacacc tctctcgnn cacgctgacc cctnctgct cttctttctac ctttcaagtt 120
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
accctgacaa gctctgagac caaatctccg atgcngtctt cgacgcttgc ctgagcagg 240
accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300
gagaa 305

<210> 2182
<211> 277
<212> nucleic acid
<213> Glycine max
<400> 2182

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cacaaagcgg gtcactgtct 60
gttcaagcta ccatctctct ctctctttct tagtgcttcc ttgccagaag ttaaaatggc 120
ccaagaaact ttcctattca catctgaatc agtgaacgag gggcacctg acaagctctg 180
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tgctgtgaa acctgcacca agaccaacat ggtgatg 277

<210> 2183
<211> 187
<212> nucleic acid
<213> Glycine max
<400> 2183

tagtgcttcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60

agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
 atgcttggag caggaccctg acagcaaggt tgcctgtgaa acctgcacca agaccaacat 180
 ggtgatg 187

<210> 2184
 <211> 282
 <212> nucleic acid
 <213> Glycine max
 <400> 2184

tcgcangcac gcgtacgtaa gctcggaatt cngctcgagc aacagcacia agcggggttac 60
 tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaaa 120
 atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180
 ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240
 aaggttgctt gtgaaacctg caccaagacc aacatggtga tg 282

<210> 2185
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 2185

gtgcangca cgcgtacgta agctcggnat tcggctcgan ctcgagccga attcgggctc 60
 gantatacaa cagcaciaaag cgggactact gtctgttcaa gactaccatc tctntctctc 120
 tttcttagtg cctccttgcc agaagttaaa atggcccaan aaactttcct attcacatct 180
 gaatcngtga acgaggggca ccctgacaag ctctgtgacc agatctccga tgctgtgctc 240
 gatgcatgct tggagcagga ccctgacagc aaggttgctt gtgaaacctg caccaagacc 300
 aacatggtga tggtt 315

<210> 2186
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 2186

anacgcangc acgcgtacgt aagctcgga ttcngctcga gggcaagccc cactcaacca 60

ccacacctct cctcggtcac gctacccctt tctgctcttc ttctaccttt caagttttaa 120
aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180
tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgctcgc agcaggaccc 240
agacagcaaa gttgctgcg aaacatgnac caaaaccaac ttggtcatgg tcttcggaga 300
aat 303

<210> 2187
<211> 297
<212> nucleic acid
<213> Glycine max
<400> 2187

acgtgcgacg cagcggtacg taagctcgga attcngctcg aggccccact caaccaccac 60
accnctcttc gttcacgcta cccctttctg ctcttcttct acctttcaag ttttaaaagt 120
ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg acaccctgac 180
aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca ggaccagac 240
agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatgggtctt cggagaa 297

<210> 2188
<211> 276
<212> nucleic acid
<213> Glycine max
<400> 2188

cgcntgcacg cgtacgttag ctcggaattc ggctcgaggc acaaagcggg ttactgtctg 60
ttcaagctac catctctctc tctctttctt agtgccctct tgccagaagt taaaatggcc 120
caagaaactt tctattcac atctgancca gtgaacgagg ggcaccctga caagctctgt 180
gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga cagcaagggt 240
gcctgtgaaa cctgcaccaa gaccaacatg gtgatg 276

<210> 2189
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2189

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 ccacacctct cctcggtcac gctacccctt tctgctcttc ttctaccttt caagttttta 120
 aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180
 tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctcg agcaggaccc 240
 agacagcaaa gttgctcgcg aaacatgcac caaaaccaac ttggtcatgg tcttcggaga 300

<210> 2190
 <211> 283
 <212> nucleic acid
 <213> Glycine max
 <400> 2190

gentangtac gcnnacgtaa gctcggaatt cggtcgcagc caccacacct ctctcgttc 60
 acgtacccc tttcngctct tcttctacct ttcaagtttt aaaagtataa agatggcaga 120
 gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 180
 aatctccgat gctgtcctcg acgcttgccct cgagcaggac ccagacagca aagttgcctg 240
 cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaa 283

<210> 2191
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 2191

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 caccacacct ctctcgttc acgtacccc tttctgctct tcttctacct ttcaagtttt 120
 aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct cgagcaggac 240
 ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 300
 gaa 303

<210> 2192
 <211> 320
 <212> nucleic acid
 <213> Glycine max

<400> 2192

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ctacctttca agtttttaaaa gtataaagat ggcagagaca ttcttattta cctcagagtc 180
ggtgaacgag ggacaccctg acaagctctg cgaccaaato tccgatgctg tctcgcacgc 240
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ggtcatggtc ttcgagaaaa 320

<210> 2193

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2193

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agtataaaga tggcagagac attcctattt acctcagagt cggatgaacga gggacaccct 180
gacaagctct gcgaccaaat ctccgatgct gtcctcgacg cttgcctcga gcaggaccca 240
gacagcaaag ttgcctgcga aacatgcacc aaaaccaact tggatcatggc ttcgagaaaa 300
a 301

<210> 2194

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2194

gcatgcacgc gtacgtaagc tcggaattcg gctcgagcca agccccactc aaccaccaca 60
ccactctctc tgctcttctt ctacctttca agtttttaaa gtattaagat ggcagagaca 120
ttcttattta cctcagagtc agtgaacgag ggacaccctg acaagctctg cgaccaaato 180
tccgatgctg tctcgcacgc ttgccttgaa caggaccag acagcaaggc tgctgcgaa 240
acatgcacca agaccaactt ggtcatggtc tcgagagat cacc 284

<210> 2195

<211> 288
<212> nucleic acid
<213> Glycine max

<400> 2195

ncacgtcgca ngcacgcnta cgtaagctcg gaattcggct cgagcaacag caciaagcgg 60
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ttaaaatggc ccaaganact ttctattca catctgaatc agtgaacgag gggcaccctg 180
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg 240
acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgg 288

<210> 2196
<211> 292
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<400> 2196

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ttcttngct tgagaaatgg caaaaaacc ttctattca catctgaatc tgtaaacgan 180
ggtcaccccg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240
caggaccctg acagcaangt tgctgtgag acatgcacca ngaccaacat gg 292

<210> 2197
<211> 316
<212> nucleic acid
<213> Glycine max

<400> 2197

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actcaaccac cacacctctc ctctttcacg ctaccctttt ctgctcttct tctacctttc 120
aagttttaaa agtataaaga tggcaganac attcctatct acctcagagt cggatgaacga 180
gggacaccct gacaagctct gcgaccaaatt ctccgatgct gtctcgacg cttgcntcga 240
gcaggacca gacagcaaag ttgctgcna nacatgcacc aaaaccaact tggatcatgg 300
cttcggagaa atcacg 316

<210> 2198
 <211> 305
 <212> nucleic acid
 <213> Glycine max

 <400> 2198

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 taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180
 ccctgacaag ctctgcgacc aaatctccga tgctgtcttc gacgcttgcc tcgagcagga 240
 ccagacagc aaagttgctt gcgaaacatg caccaanacc aacttggtca tggctctcgg 300
 agaaa 305

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<210> 2199
 <211> 301
 <212> nucleic acid
 <213> Glycine max

 <400> 2199

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 acgctctgct tccagcgagt gttctttctt cgtttcaaca ccttaatttg cacacgctgc 120
 ttcttcagct tgagaaatgg cacaagaaac ctttctattc anatctgaat ctgtaaacga 180
 gggtcacccc gacaagctgt gngaccagat ctctgatgca gtgcccgatg cgtgccttga 240
 acaggncctt gacancaagg ttgcctgtga gacatgnacc aagaccaana tggatcatgtt 300
 t 301

<210> 2200
 <211> 289
 <212> nucleic acid
 <213> Glycine max

 <400> 2200

 gtcgcangca cgcgtacgta agctcggaa ttcggctcga gacttaacaa cagcacaag 60
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120
 aagttaaaat ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180

ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc 240
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccna catggtgat 289

<210> 2201
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 2201

gtcgcattgca cgcctacgta agctcggaat tnnctcagag gccaggcaa gcccactca 60
 accaccacac ctctctctgt tcacgctacc cttttctggc tcttcttcta cttttcaagt 120
 tttaaaagta taaagatggc agagacattc ctatttacct cagatcggt gaacgaggga 180
 caccctgaca agctctgcca ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240
 gaccagaca gcaaagttgc ctgcgaaaca tgcaccanaa ccaacttggt catggtcttc 300
 ggagaaatc 309

<210> 2202
 <211> 250
 <212> nucleic acid
 <213> Glycine max
 <400> 2202

gcagacttaa caacagcaca aagcgggtta ctgtctgttc aagctaccat ctctctctct 60
 nctttcttag tgctctcttg ccagaagtta aaatggccca agaaactttc ctattcacat 120
 ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180
 tcgatgcatg cttggagcag gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaaga 240
 ccaacatggt 250

<210> 2203
 <211> 295
 <212> nucleic acid
 <213> Glycine max
 <400> 2203

gtcgcattgca cgcgtacgta agctcggaat tcggctcagag gcccactca accaccacac 60
 ntctctctgt tcacgctacc cttttctgct cttcttctac ctttcaagtt ncaaaagtat 120

aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240
caaagttgcc tgcgaaacat gcaccaaaac caacttggtc atggtcttcg gagag 295

<210> 2204
<211> 272
<212> nucleic acid
<213> Glycine max
<400> 2204

gncgengcac gcgtacgtna gctcgggnatt cggctcgagg gcccactca accaccacac 60
cactctctct gctcttcttc tacctttcaa gtttttaaag tattaagatg gcagagacat 120
tctattttac ctcagagtca gtgaacgagg gacacctga caagctctgc gaccaaactc 180
ccgatgctgt cctcgacgct tgccttgaac aggaccaga cagcaagggt gcctgcgaaa 240
catgcaccaa gaccaacttg gtcatggtct tc 272

<210> 2205
<211> 276
<212> nucleic acid
<213> Glycine max
<400> 2205

cgtcgcangc acgcgtacgt aagctcgga ttcggctcga gccaaagccc actcaaccac 60
cacaccactc tctctgctct tcttctacct ttcaagtttt taaagtatta agatggcaga 120
gacattccta ttacctcag agtcagtga cgagggacac cctgacaagc tctgcgacca 180
aatctccgat gctgtctctg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240
cgaaacatgc accaagacca acttggtcat ggtctt 276

<210> 2206
<211> 307
<212> nucleic acid
<213> Glycine max
<400> 2206

ntontatgca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gcccactca 60
accaccacac ctctctctgt tcacgtacc cttttctgct cttctcttac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240
 acccagacag caaagttgcc tgcgaaacat gcacccaaac caacttggtc atggtcttcg 300
 gagaaat 307

<210> 2207
 <211> 311
 <212> nucleic acid
 <213> Glycine max
 <400> 2207

tcgcangcac gcgtagctna gctcggaatt cggctcgagg ccatttggn agttaggttc 60
 tgcaagctct gcttcacgag agtggtcttt ctctgtttca acaccttaat ttgcacacgc 120
 tgcttcttca gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180
 cgaggggtcac cccgacaagc tgtgcgacca gatctctgat gcagtgcctg atgcgtgcct 240
 gaacaggacc ctgacagcaa ggttgctgt gagacatgca ccaagaccaa catggtcagg 300
 tcttgagag a 311

<210> 2208
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2208

nnnccatgc acgcgtacgt aagctcgga ttcggctcga ggactgttat gtttaaattg 60
 tagtcatggt ggtgtttttg gctgtgaatt tgctcatatg tgctaattat gtgttcttgt 120
 ttgatgttac tctacagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 180
 agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctnccatg 240
 catgcttggg gcaggaccct gacagcaang ttgcctgtga aacctgcacc aagaccaaca 300
 tggatgatgt 310

<210> 2209
 <211> 338
 <212> nucleic acid
 <213> Glycine max

<400> 2209

tcgcatgcac gcgtacgtaa gctcggaatt cnnctcgagg caagccccac tcaaccacca 60
cacctctect cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacaccctga 180
caagctctgc gaccaaactc cgatgctgtc ctcgacgctt gcctcgagca ggaccagac 240
agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatggtctt cggagaaatc 300
acgaccaggc caagttgatt acgagaagta gtgcgtga 338

<210> 2210

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2210

antencangc acgcgtacgt aagctcgga ttcggctcga gaacagcaca aagcgggtta 60
ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120
aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180
gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttgganccag gaccctgaca 240
gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggt gatggttt 288

<210> 2211

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2211

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gccccactca 60
accaccacac ctctcctcgt tcacgctacc ctttctgctc ttcttctacc tttcaagttt 120
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 300
gaaatcacga c 311

<210> 2212

<211> 328
 <212> nucleic acid
 <213> Glycine max

 <400> 2212

 angtcctcang cacgcgtacg taagctcgga attcagctcg agcngctcga gcacacctct 60
 cctcggttcac gctacccctt tctgctctnc ttctaccttt caagttttna angtnataag 120
 gtggcagaga cattcctatt tacctcagag tcgntgaacg agggacaccc tgnnaagctc 180
 tgcgaccaa tctccgatgc tgtcctcgac gcttgccctcg agcaggaccc agacagnaaa 240
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 agccaacgt tgatacgaga agatatgc 328

<210> 2213
 <211> 309
 <212> nucleic acid
 <213> Glycine max

 <400> 2213

 acgtcgcang cacgcgtacg taagctcgga attcggctcg agcaagcccc actcaaccac 60
 cacacctctc ctggttcacg ctaccccttt ctgctcttct tctacctttc aagttttaaa 120
 agtataaaga tggcagagac attcctatct acctcagagt cggatgaacga gggacaccct 180
 gacaagctct ggcaccaaatt ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240
 gacagcaaag ttgcctgcga aacatgcacc aaaaccaatt ggtcatgggtc ttcggagaaa 300
 tcacgacca 309

<210> 2214
 <211> 299
 <212> nucleic acid
 <213> Glycine max

 <400> 2214

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 tcgacttcag acctggaatg atcaccatta acttgacct taagaggggt ggtcataggt 120
 tcctcaagac agctgcttat ggacactttg gaaggatga tgcagacttc acctgggaag 180
 ttgtgaagcc actcaagtca gagaagcctc aagcttaaga gtgttggtta gttaatcact 240

cccttcagtg gatgtcttgc tgggtgtgga tgaataattt gcgtgtttca tgactacta 299

<210> 2215
 <211> 297
 <212> nucleic acid
 <213> Glycine max

<400> 2215

ngtgcangc acgcgtacgt aagctcggaa ttcggctcga ggccaggcaa gcccactca 60

accaccacac ctctctctgt tcacgtacc cttttctgct cttctttctac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240

accagacag caaagttgcc tgcgaaacat gcacccaaac caacttggtc atggtct 297

<210> 2216
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 2216

gtgcangca cgcgtacgtn agctcggaat tcggctcggg ggccaggcaa gcccactca 60

accaccacac ctctctctgt tcacgtacc cttttctgct cttctttctac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240

accagacag caaagttgcc tgcgaaacat gcaccannac caacttggtc atggtctt 298

<210> 2217
 <211> 284
 <212> nucleic acid
 <213> Glycine max

<400> 2217

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gttactgtct gttcaagcta ccattctctc ctctctttct tagtgctcc ttgccagaag 120

ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcaccctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggaccctg 240

acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtg 284

<210> 2218
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2218

ttgcatgcan gcgtacgtaa nctcggaat tcggctcgag nggagttagg ttctgcacgc 60
 tctncttcca gcgagtgttc tttctctgtt tcaacacctt aatttgcang acgtgcttn 120
 tnaaganttg agaaatggca caagaaacct ttctattcac atctgaatct ntaaagcagg 180
 gtcaccccgga naagctgtgc gancagatct ctnatgcagt gctcgatgcg tgccttgaac 240
 aggaacctga cagcaaggtt gcctgtgaga catgcaccaa gaccaacatg gcatggtc 298

2219

<210> 2219
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 2219

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 aaccaccaca cctctcctcg ttcacgtac cctttctgc tcttcttcta cctttcaagt 120
 tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 180
 caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240
 gaccagaca gcaaagttgc ctgcgaaaca tgcagcaaaa ccaacttggc catggtcttc 300
 ggn 303

<210> 2220
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 2220

ttacatgcac acgtacgtaa gctcggaatt cngctcgaga ggcaagcncc actcaaccac 60
 cacacctctc ctggttcacg ctaccccttt ctgctcttct tctacctttc nagttttaaa 120
 agtataaaga tggcagagac attcctatctt acctcagagt cggatgaacga gggacacct 180

gacaagctct gcgaccaaact ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240
gatagcaaag ttgcctgcna aacatgcacc aaaaccaact tggatcatggt cttcggagaa 300
a 301

<210> 2221
<211> 304
<212> nucleic acid
<213> Glycine max
<400> 2221

agtcgcangc acgcgtacgt aagctcggaa ttcggtctga ggggccaggc aagccccact 60
caaccaccac acctctctctc gttcacgcta cccctttctg ctcttcttct acctttcaag 120
ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240
ggaccagac agcaaagttg cctgcgaaac atgcaccaga accaacttgg tcatggtctt 300
cgga 304

<210> 2222
<211> 311
<212> nucleic acid
<213> Glycine max
<400> 2222

tancgcatgc acgcgngtaa nntnnnaatc ggnattcggc tcgagtttga ggccaggcaa 60
gccccactca accaccacac ctctctctctg tcacgctacc cttttctgct cttcttctac 120
ctttcaagtt ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg 180
aacgagggac accctgacaa gctctgcgat caaatctccg atgctgtcct cgacgcttgc 240
ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacaaaac caacttggtc 300
atggtcttcg g 311

<210> 2223
<211> 284
<212> nucleic acid
<213> Glycine max
<400> 2223

annctaattgc acgcgtacgt aagctcggaa ttccggctcga gaacaacagc acaaagcggg 60
 ttactgtctg ttcaagctac catctctctc tctctttctt agtgccctct tgccagaagt 120
 taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg ggcaccctga 180
 caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga 240
 cagcaagggtt gcctgtgaaa cctgcaccaa gaccaacatg gtga 284

<210> 2224
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 2224

nncnaannnn cgcattgcacg cgtacgtaag ctccggaattc ggctcgaggc agacttaaca 60
 acagcacaaa gcgggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc 120
 ctcttgcca gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga 180
 cgaggggac cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcattgctt 240
 ggagcaggac cctgacagca aggttgctg tgaaacctgc accaagacca acatggtga 299

<210> 2225
 <211> 324
 <212> nucleic acid
 <213> Glycine max
 <400> 2225

agtcgcangc acgcgtacgt aagctcggaa ttccggctcga gatttgaggc caggcaagcc 60
 ccactcaacc accacacntc tncncgttca cgtaccctct ttctgnctct tcttgctncc 120
 tttcaagttt taaaagtata aagatggcag agacattcct atttacctca gagtcgggtga 180
 acgagggaca cctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240
 tcgagcagga ccagacagc aaagttgcct gcganacatg caccaaaacc aacttggtca 300
 tggctctngga gaaatcacga ccaa 324

<210> 2226
 <211> 304
 <212> nucleic acid
 <213> Glycine max

<400> 2226

ntcgcangca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gcccactca 60

acnaccacac ctctctctgt tcacgctacc cttttctgct cttcttctac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgcgac caaatctccg atgctgtcct cgncgcttgc ctcgagcagg 240

accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300

gaga 304

<210> 2227

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2227

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc cactcaacc 60

accacacctc tctctgttca cgtaccctt ttctgtcttt cttctacctt tcaagtttta 120

aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180

ctgacaagct ctgcgaccaa atctccgatg ctgtctctga cgcttgcttc gagcaggacc 240

cagacagcaa agttgcttgc gaaacatgca gcaaaaccaa cttggtcatg gtcttcggag 300

<210> 2228

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2228

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gggccaggca agccccactc 60

aaccaccaca cctctctctg ttcacgctac ccttttctgc tcttcttcta cctttcaagt 120

tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180

cacctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240

gaccagaca gnaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt catggtcttc 300

gg 302

<210> 2229

<211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2229

 gtcgcangca cgcgtacgta agctcggaat tcggctcgag caggcaagcc ccaactcaacc 60
 accacacctc tctctgttca cgtaccctt ttctgctctt cttctacctt tcaagtttta 120
 aaagtataaa gatggcagag acatttctat ttacctcaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240
 cagacagcaa agttgcctgc gaaacatgca ncaaaaccaa cttggtcatg gtcttcgg 298

<210> 2230
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2230

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 ccaccacacc tctctcggtt cagctaccc ctttctgctc ttcttctacc tttcaagttt 120
 taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240
 ccagacagc aaagttgcct gcgaaacatg caccaaaacc aacttggtca tggctcttc 298

<210> 2231
 <211> 269
 <212> nucleic acid
 <213> Glycine max

 <400> 2231

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 tttctnctct tcttctacct ttcangtttt aaaagtataa agatggcaga gacattccta 120
 tttacctcag agtcggtgaa cgaggacac cctgacaagc tctgcgacca aatctccgat 180
 gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 240
 accaaaacca acttggtcat ggtcttcgg 269

<210> 2232

<211> 290
 <212> nucleic acid
 <213> Glycine max

 <400> 2232

 ncaattaana gtcgcangca cgcgtacgta agctcggaat tcggctcgag gttaggttct 60
 gcacgctctg ctccacgca gtgttctttc ttcgtttcaa caccttaatt tgcacacgct 120
 gcttcttcag cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac 180
 gagggtcacc ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt 240
 gaacaggacc ctgacagcaa ggttgccctgt gagacatgca ccaagaccaa 290

<210> 2233
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 2233

 gtcgcangca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gccccactca 60
 accaccacac ctctctctgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
 accctgacaa gctctgacgac caaatctccg atgctgtcct cgacgcttgc ctgagcagg 240
 accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300
 gagaaa 306

<210> 2234
 <211> 311
 <212> nucleic acid
 <213> Glycine max

 <400> 2234

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 accaccacac acgctctctg tncacgctac cctttctggt ctcttcttct accttcaag 120
 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240
 ggaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt catggtcttc 300

ggagaaatca c

311

<210> 2235
<211> 289
<212> nucleic acid
<213> Glycine max

<400> 2235

natcgcacatgc acgcgtacgt nagctcggaa ttccggctcga gcaacagcac aaagcggggtt 60
actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg ccagaagtta 120
aatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctganc 180
aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240
agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatggttt 289

<210> 2236
<211> 260
<212> nucleic acid
<213> Glycine max

<400> 2236

agcagactta acaacagcac aaagcggggtt actgtctgtt caagctacca nnnnnnnnnn 60
nnnnnnntag tgcctccttg ccagaagtta aatggccca agaaactttc ctattcacat 120
ctgaatcagt gaacgagggg caccctncac aagctctgtg accagatctc cgatgctgtg 180
ctcgatgcat gcttggagca ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag 240
accaacaatgg tgatggttt 260

<210> 2237
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 2237

ntcanntacg cgtangtanc actgcgtacn tnagctcggaa attcggctcg agcagcacia 60
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120
agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180
ccctgtacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atgggtttt 298

<210> 2238
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 2238

tcgcngaacg ngtagcgttaag ctgcgaattc ggctcgangn catttgggag ttaggtttna 60
 acgctcngcg tnnagngagt gatntttctt cgtntcanca cntnaaattg cancacgctg 120
 cttcttcngc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180
 anggtcaccg cgacaagctg tgtgaccaga tctctgatgc antgctcgat gcgngccttg 240
 aacaggaccc tgacagcaag ttgctgtga gacatgcacc atgaccaaca tggtcaggtc 300
 n 301

<210> 2239
 <211> 309
 <212> nucleic acid
 <213> Glycine max
 <400> 2239

acgtcgcang cagcgttacg taagctcgga attcngctcg aggcaagccc cactcaacca 60
 ccacacctct cctcgttcac gctaccctt tctgctactt cttctacctt tcaagtttta 120
 aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgcttc gagcaggacc 240
 cagacagcaa agttgcctgc gaacatgcac caaaaccaac ttggtcatgg tcttcggaga 300
 aatcacgac 309

<210> 2240
 <211> 293
 <212> nucleic acid
 <213> Glycine max
 <400> 2240

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gttaacaaca gcacaaagcg 60
 ggttactgtc tgttcangct accatctctc tctctctttc ttagtgcttc cttgccagaa 120

gttaaaatgg cccaagaaac ttctctattc acatctgaat cagtgaacga ggggaccctg 180
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggncctg 240
acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttc 293

<210> 2241
<211> 279
<212> nucleic acid
<213> Glycine max

<400> 2241

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tgtgttcaag ctaccatctc tctctctctt tcttagtgcc tcttgccag aagttaaaat 120
ggcccaagaa actttctctat tcacatctga atcagtgaac gaggggcacc ctgacaagct 180
ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240
ggttgctctg gaacctgcac caagaccaac atgggtgatg 279

<210> 2242
<211> 181
<212> nucleic acid
<213> Glycine max

<400> 2242

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agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
g 181

<210> 2243
<211> 289
<212> nucleic acid
<213> Glycine max

<400> 2243

acgtcgcang cacgcgtacg taagctcggg attcggctcg aggcagactt aacaacagca 60
caaagcgggt tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt 120
gccagaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg 180

2244
 287
 nucleic acid
 Glycine max
 2244
 60
 120
 180
 240
 287
 2245
 310
 nucleic acid
 Glycine max
 2245
 60
 120
 180
 240
 300
 310
 2246
 284
 nucleic acid
 Glycine max
 2246
 60
 120

gcaccctgac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca 240

ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaacaatg 289

<210> 2244

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2244

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aagogggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120

cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180

accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatg 287

<210> 2245

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 2245

aagtcgcatg cagcgtacg naagctcggc attnggcncg cggncaggc aagcaccact 60

caaccancac acatctnctc gttcaagcta cccctttgtn cncncttctt aantttcaag 120

ttttaaaagt atacagatgg cagagacatt cctatttacc tcagagtggg tgaacgaggg 180

acacntgac aagctctgcg accaaatctc cgatgctgtc ctcgacgott gcntcgagca 240

ggaccagac agcaangttg cctgcgaaac atgcacnga accaacttgg tcatggtctt 300

cggagaaatc 310

<210> 2246

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2246

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tcacgctacc cttttctgct cttcttctac ctttcaagtt tnaaangtnt aaagatggca 120

gagacatncc tatttacctc agagtcggtg aacgagggac acccngacaa gctctgcgac 180
 caaanctccg atgcngtcct cgacgcttgc ctcgagcagg acccagacag caaagntgcc 240
 tgcgaaacan gcacaaaaac caacttggtc atggtcttcg gaga 284

<210> 2247
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 2247

ntntcgcan gacgcgtacg taagctcgga attcggtcag aggccaggca agccccactc 60
 aaccaccaca ctggctcctc gttnacgcta cccctttctn cctctttctc tacctttcaa 120
 gttttaaaag tataaagatg gcagagacat tctatattac ctgagagtcg gtgaacgagg 180
 gacaccctga caagctctgc gaccaaattc ccgatgctgt cctcgacgct tgccctgagc 240
 aggaccaga cagcaaagtt gcttgcgaaa catgcaccaa aaccaacttg gtcattggtc 299

<210> 2248
 <211> 182
 <212> nucleic acid
 <213> Glycine max
 <400> 2248

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 agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
 atgcttgag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180
 gg 182

<210> 2249
 <211> 313
 <212> nucleic acid
 <213> Glycine max
 <400> 2249

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 ancnggtta cttnctgtnc aagctancca tctctctctc tctttcttag tgctccttg 120
 ccagaagtta aaatggccca agaaactttc ctannacat ctgaatcagt gaancgagg 180

gcacctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcneg nttggagcag 240
gaccctgaca gcaaggttgc ctgtgaaacc ngcaccaaga ccaacatggt gatgggttttc 300
ggaganntca caa 313

<210> 2250
<211> 289
<212> nucleic acid
<213> Glycine max
<400> 2250

gcacgcgtac gtaagctcgg aattcggctc gagcaggcaa gcccactca accaccacac 60
ctctcctcgt tcacgctacc cttttctgct cttctttctac ctttcaagtt ttaaaagtat 120
aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240
caaagttgcc tgcgaaacat gcancaaaac caacttggtc atggtcttc 289

<210> 2251
<211> 264
<212> nucleic acid
<213> Glycine max
<400> 2251

atttgaggcc aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt 60
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tacctcagag tcggtgaacg agggacaccc tgacaagctc tgcgaccaaa tctccgatgc 180
tgtcctcgac gcttgctcgc agcaggaccc agacagcaaa gttgcctgcg aaacatgcac 240
caaaaccanc ttggtcatgg tctt 264

<210> 2252
<211> 315
<212> nucleic acid
<213> Glycine max
<400> 2252

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acacctctcc tcgttcacgc tacccttttc tgctctttct ctacctttca agtttttaaa 120

gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 180
 aaagctctgc gaccaaattct ccgatgctgt cctcgacgct tgcctcgagc aggaccaga 240
 cagcaaagtt gcctgcgaaa catgcacnaa aaccaattgg tcatgggtctt cggagaaatc 300
 acgaccaagg ccaag 315

<210> 2253
 <211> 191
 <212> nucleic acid
 <213> Glycine max
 <400> 2253

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 agtgaacgag gggcaccctg acaagcttgt gaccagatct ccgatgctgt gctcgatgca 120
 tgcttggagc aggaccctga cagcaaggtt gcctgtgaaa cctgcaccaa gaccaacatg 180
 gtgatggttt t 191

<210> 2254
 <211> 304
 <212> nucleic acid
 <213> Glycine max
 <400> 2254

gttgcangca cgcgtacgta agctcggaat tcggctcgag agcagactta acaacagcac 60
 aaagcgggtt actgtctgtt caagctnca nctctctctc tctttcttag tgctccttg 120
 ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacganggg 180
 caccctgtgc aagctctgtg accagatctc cgatgctgtg ctcgatgcat gattggagca 240
 ggaccctgac agcaaggttg nctgtgaaac ctgcaccaag ancaacatgg tgatggtttt 300
 cgga 304

<210> 2255
 <211> 317
 <212> nucleic acid
 <213> Glycine max
 <400> 2255

tcgcatgcnc gcgtacgtna gctcgggnatt cggctcgagc anaangcngg ttactgtctg 60

ttcaagctac catctctctc tctctttctt antgcctcct tgccagangt taaaatggcn 120
 caagaanctt tcttattcac atctgaatnn gtgaacgagg ggcaccctga acaanctctg 180
 tgancagatc tccgatgctg tgctcgntgc atncttgag caggaccctg acagcnaggt 240
 tncctgtgna acntgcacca agnccancat ggngatgggt ttcggagann tcacaaccan 300
 ggccaacgtg gactatg 317

<210> 2256
 <211> 235
 <212> nucleic acid
 <213> Glycine max
 <400> 2256

cngcacgctg acgtaagctc ggaattcggc tcgagggaga ttggtgctgg tgaccaaggt 60
 catatgttgc gctatgnnet gacgangntc ccgagctcat gcccntgagc catgtccttg 120
 ccacgaagct cgggtgtcaag ctcanagagg ttcggaanaa cgggacatgc ccttggtgta 180
 ganctgntgg caagaccnag gtcantgttg nnnactacaa tggcaagggn gccat 235

<210> 2257
 <211> 319
 <212> nucleic acid
 <213> Glycine max
 <400> 2257

cgatgcacgc gtacgtaagc tcggaattcg gctcgagttt ggggagttag gttctgcacg 60
 ctctgcttcc agcgagtgtt ctttcttcgt ttcaacaacc ttaatttgca cacgctgctt 120
 cttcagcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180
 gtcaccccca caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240
 aggaccctga cagcaagttg cctgtgagac atgcaccaag accaactggt caggtctttg 300
 gagagatcac aaccagggc 319

<210> 2258
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2258

gtcgcacgca cgcgtacgta agctcggaat tccggtcgag caggcaagcc cactcaacc 60
accacacctc tctcggttca cgtacacctt tctgctctt cttctacctt tcaagtttta 120
aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgacct gagcaggacc 240
cagacagcaa agttgacctg gaaacatgca acaaaaccaa ttggtcatgg tcttcggaga 300
aatcac 306

<210> 2259
<211> 300
<212> nucleic acid
<213> Glycine max

<400> 2259
acgtcgang cagcgtacg taagctcgga attcggtcg agcttaacaa cagcacaag 60
cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tcttgccag 120
aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggacacc 180
ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240
ctgacagcaa ggttgacctg gaaacctgca accaagacca acatggtgat ggnttncgga 300

<210> 2260
<211> 330
<212> nucleic acid
<213> Glycine max

<400> 2260
gtcgcacgca cgcgtacgta agctcggaat tnggctcgag ctcgagccgc aggaaagccc 60
cactcaacca ccacacctct cctcggttac gctacacctt aactgcttct tcttctacct 120
ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa 180
cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgacct 240
cgagcaggac ccagacagca aagttgacct cgaaacatgc accaaaacca cttggtcatg 300
gtcttcggag aaatcagacc aaggccaagt 330

<210> 2261
<211> 180
<212> nucleic acid

<213> Glycine max

<400> 2261

tagtgccctcc ttgccagaag ttaaaatggc ccaagaaact ntcctattca catctggatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcnntgc 120
atgcttggag caggancctg acagcaaggt tgccctgtgaa acctgcacca agaccaacat 180

<210> 2262

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2262

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag caagccccac tcaaccacca 60
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctacagatcg gtgaacgagg gacaccctga 180
caagctctgc gaccaaactc ccgatgctgt cctcgacgct tgccctcgagc aggaccacga 240
cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatg 286

<210> 2263

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2263

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tttgaggcca ggcaagcccc 60
actcaaccac cacacctctc ctgcgttcacg ctaccctttt ctgctcttct tctacctttc 120
aagtttttaa agtataaaga tggcagagac attcctatth acctcagagt cggatgaacga 180
gggacaccct gacaagctct gcgaccaaact ctccgatgct gtccctcgacg cttgcctcga 240
gcaggaccca gacagcaaag ttgcctgcga aacatgcacc aaaaccnact tggatcatggt 300

<210> 2264

<211> 332

<212> nucleic acid

<213> Glycine max

<400> 2264

cgcangcacg cgtacgtaag ctcggaattc ggctcgagcg acttaacaac agcacaaagc 60
 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct cnttgccaga 120
 agttaaaatg gccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgnt gcatgcttgg agcaggaccc 240
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atgtgatggg ttccggagagn 300
 tcacaaccan gcaacgtgga ctatgagagg tt 332

<210> 2265
 <211> 274
 <212> nucleic acid
 <213> Glycine max

<400> 2265
 tcacnggtac gtnagctcgg aattcggctc nagcttaaca acagcacaaa gcgggttact 60
 gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctccctgcca gaagttaaaa 120
 tggoccaaaga aactttccta ttcacatctg aatcagtga cggaggggcac cctgacaagc 180
 tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240
 aggttgcttg tgaaacctgc accaagacca acat 274

<210> 2266
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 2266
 gcnannnagg nnagtcgcan gcacgcgtac gtaagctcgg aattcggctc gaggcagact 60
 taacaacagc acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt 120
 agtgctcctt tgccagaagt taaaatggcc caagaaactt tcctattcac atctgaatca 180
 gtgaacgagg ggcaccctga caagctctgt gaccagatct ccgatgctgt gctcgatgca 240
 tgcttgagc aggaccctga cagcaagggt gcctgtgaaa cctgcaccaa gaccaacatn 300

<210> 2267
 <211> 288
 <212> nucleic acid
 <213> Glycine max

<400> 2267

ngtcgcangc acgcgtacgt aagctcggaa ttcngctcga ggccccactc aaccaccaca 60
cntctcctcg ttcacgctac ccttttctgc tcttcttcta cctttcaagt tttaaaagta 120
taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca 180
agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
gnaagttgc ctgcgaaaca tgcacaaaaa ccaacttggc catggtct 288

<210> 2268

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2268

gttgcangca cgcgtacgta agctcggaa ttcggtcgcg atttgaggtt tggagcgact 60
naactaatca ttaatttgca ctgcgtgttt cagcttcac accctttctt ttgcatcatt 120
tatatctctt gagaaatggc acaagaaacc tntctattca catctgaatc tgtaaacgag 180
ggtcaccccg acangctgtg cgancagatc tctgatgcag tacttgatgc gtgccttgaa 240
caggaccctg acagcaaggt tgcngtgag acatgnacca agaccaacat g 291

<210> 2269

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2269

cacgcgtacg taagctcgga attcggctcg agncaaagga gtgatttgga gtttgagcg 60
actgaactaa tcattaattt gcactcgtg tttcagcttc atcacccttc ttttgcata 120
tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180
agggtcaccc cgacangctg tncnaccaga tctctgatgc agtacttgat gcgtgccttg 240
aacnggaccc tggacagcaa ggttgctgt gagacatgca ccaagaccaa catggtct 298

<210> 2270

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2270

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60

aagcgggtta ctgtctgttc aagctaccat ctncctctctc tctttcttag tgcctccttg 120

ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180

caccctgaac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca 240

ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatgg 296

<210> 2271

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2271

gtcncatgca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcacaaaag 60

cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120

aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180

ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240

nncgacagca aggttgccctg tgaaacctgc accaagacca acatgggtg 288

<210> 2272

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2272

nagtcgcatg cagcgtacg taagctcgga attcngctcg aggcaagccc cactcaacca 60

ccacacctct cctcgttcac gctaccctt tctgctcttc ttctaccttt caagttttta 120

aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180

tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctg agcaggaccc 240

agacagcaaa gttgcctgcg aaacatggca ccaaaaccaa cttggtcatg gtcttcgga 299

<210> 2273

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2273

gcngcacgcg tacgtaagct cggaattcgg ctcgaggcca ggcaagcccc actcaaccac 60
cacacctctc ctcgttcacg ctaccccttt ctgctcttct tctacctttc aagttttaaa 120
agtataaaga tggcagagac attcctatctt acctcagagt cggatgaacga gggacaccct 180
gacaagctct gcgaccaaatt ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240
gcagaccaaag ttgctcgcga aacatgcacc anaaaccaac ttggtcatgg tcttcggaga 300

<210> 2274

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 2274

acgtcgcang caogcgtagc taagctcggg attcggctcg agggagtttg gagcgactga 60
actaatcatt aatttgcact cgctgtttca gcttcacac ctttcttttg catcatttat 120
atctcttgag aaatggcaca agaaaccttt ctattcacat ctgaatcgta aacnagggtc 180
accccgacaa gctgtncnat cagatctctg atgcagtact tgatgcntgc cttgancagg 240
nccctgacag caaggttgcc tgtgagacnt gcaccaagac caacatgggc atggtct 297

<210> 2275

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2275

gcangcacgc gtacgtnagc tcggaattcn gctcgaggcc cactcaacc accacacctc 60
tcctcgttca cgctacccct ttctgctctt cttctacctt tcaagtttta aaagtataaa 120
gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc ctgacaagct 180
ctgcgacnna atctccgatg ctgtcctcga cgcttgccct gagcaggacc cagacagcaa 240
agttgcctgc gaaacatgca ccaaaaccaa ttggtcatgg tcttcggaga aatcac 296

<210> 2276

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2276

acgtcgang cagcggtacg tnagcnnccg aattcngctc gagcaggcaa gcccactca 60

accaccacac ctctctctgt tcacgtacc cttttctgct cttttcttac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgacgac caaatctccg atgctgtcct cgacgcttgc ctgagcagg 240

accagacag caaagttgcc tgcgaaacat gcacaaaac caattggtca tggctctcgg 300

agaaat 306

<210> 2277

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2277

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagg gccaggcaag ccccactcaa 60

ccaccacacn nctctctgtt cagcgtaccc cttttctgct ttctttctacc tttcaagttt 120

taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180

ccctgacaag ctctgacgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240

cccagacagc aaagttgcct gcgaaacatg caccaaaaacc aacttgg 287

<210> 2278

<211> 206

<212> nucleic acid

<213> Glycine max

<400> 2278

gccacttacc anaaaacctg aagaaattgg tgctgntgac cagggtcaca tgtttggtca 60

tgccactgat gaaaccctg aattgatgcc attgagccat gttcttgcaa caaaactcgg 120

tgctcgtctc accgaggttc gcnagaacgg tactggcctt ggctgangct gatggaagac 180

ccaagtgacc gttgagtata caatga 206

<210> 2279

<211> 265

<212> nucleic acid

<213> Glycine max

<400> 2279

tgcacgcacg cgtacgttag ctcggaattc ggctcgagac agcacaggag cgggttaacng 60
tctgttcaag ctaccatctc tctctctctt tcntagtgcc tccttgccag aagttaaaat 120
ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagcn 180
ctgcgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240
ggttgctctg gaaacctgca ccaag 265

<210> 2280

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2280

nanngcncgn gtacgtaagc tcgganttcg gctcgagggc caggcaagcc ccaactcaacc 60
accacacctc tctcgttca cgctaccctt tngtgcctt cttctacctt tcaagtttta 120
aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
ctgacaagct ctgcgaccaa ntctccgatg ctgtcctcga cgcttgctc gagcaggacc 240
cagacagcaa agttgctgc gaaacatgca ccataaccaa cttggctatg g 291

<210> 2281

<211> 330

<212> nucleic acid

<213> Glycine max

<400> 2281

gcacgcgtac gtaagctcgg aattcggctc gagatttgag gccaggcaag cccactcaa 60
ccaccacacc tctcctcgtt cacgctaacc cttctgctc ttcttctacc tttcaagttt 120
tanaagtata aagatggcag agacattcct attacctca gagtcggtga acgagggaca 180
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc togagcagga 240
cccagacagc aaagttgcct gcgaaacatg gcacaaaac caattggtca tggctctcgg 300
agaaatcaga ccaggccaag ttgattacga 330

<210> 2282

<211> 283

<212> nucleic acid

2282
2283
2284
2285

<213> Glycine max

<400> 2282

tcgcangcac gcgtacgtaa gctcggaatt cngctcgagg caagccccac tcaaccacca 60
 cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
 tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacaccctga 180
 caagctctgc gaccaaatct ccgatgctgt cctcgacgct tgcctcgagc aggaccacaga 240
 cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtc 283

<210> 2283

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2283

ncgtcgcatg cncgcgtacg taagctcgga attcggtctg ngngccaggn aagccccact 60
 caaccaccac acctctctctc gttcacgcta cccctttctg ctctttctct acctttcang 120
 tttttaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgnacgaggg 180
 ncacctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgngcn 240
 ggaccagac agcaaagttg cctgcgaaac atgcaccaat acnaacttgg tcatggtctt 300
 cg 302

<210> 2284

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2284

nangcacgcg tacgtaagct cggaattcgg ctcgagcaga gttgagacca agacacactc 60
 gttcatatat ctctctgctc ttctcttctc ttctacctct caagtttttg aagtataaag 120
 atggcagaga cattcctatt cacctcggag tcagtgaacg agggacaccc tgataagctc 180
 tgcgacaaa tctccgatgc tgtcctcgac gcttgctcgc aacaggaccn nacaancaag 240
 gttgcctgng aaacatgcac caagaccaac ttggtcatgg tctn 284

<210> 2285

<211> 208
 <212> nucleic acid
 <213> Glycine max

<400> 2285

cangcacgcg tacgtaagct cggaattcgg ctcgagcatt ggtgtccctg agcccttgtc 60
 agtgtttgtg gacacttatg gaactgggaa gattcctgac aaggagattc tgcaaattgt 120
 gaaggagaat ttcgacttca gacctggaat gatcaccatt aacttggacc ttaanagggg 180
 tggtcatagg ttcttcaaga canntgct 208

<210> 2286
 <211> 270
 <212> nucleic acid
 <213> Glycine max

<400> 2286

gcagacttna caacagcaca naggcgggta ctgtctgttc aangctacga tctctgtctc 60
 tgttatctta gtgcctacct tgccagaagn nannatggcc caatnaantt tccnattcac 120
 atctgantca ntgaacnatg ggcaccctga naanctctgt gnccagatct ccgatgctgt 180
 gctcgatgca tgcttggagc aggaccctga nagnaggtt gcntgtnaaa cctgnaccaa 240
 gaccaacatg gtgatggtn tggagagat 270

<210> 2287
 <211> 302
 <212> nucleic acid
 <213> Glycine max

<400> 2287

nntcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggntttgagg ccaggcaagc 60
 ccactcaac caccacacnt ctctcgttc acgtacccc tttctggctc ttcttctacc 120
 tttcaagttt taaaagtata aagatggcag agacattcct atttacctca gagtcgggtga 180
 acgagggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240
 tcgagcagga ccagacagc aaagttgcct gcgaaacatg caccanaacc aacntgggtca 300
 tg 302

<210> 2288

<211> 195
 <212> nucleic acid
 <213> Glycine max

 <400> 2288

 cgtaagctcg gaattcggct cgagncaatg atcaaattgc tgcggacctt aaagagcatg 60
 ttatcaagcc tgtcattctt gagaagtacc ttgatgagaa ncaccatctt ccaccttaac 120
 ccttctggcc gttttgtcat tgggtggcct catggtgatg ctggtctcac tggaagaaaa 180
 tcatcattga tacct 195

<210> 2289
 <211> 314
 <212> nucleic acid
 <213> Glycine max

 <400> 2289

 ngcntgtacg cgtacgtaag ctcggaattc ggctcgaggt caggcaagcc ccaactcaacc 60
 accacacctc tgectgngtt cangctaccc ctttntgctc ttcttctacc tttgaagttt 120
 taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acganggaca 180
 cctgacaag ctctgcganc caaatctccg atgctgtcct cgacgnittgn ctcgagnagg 240
 acccagacag naaagttgcc tgcgatanat gcaccannac caacttggtc atggtcttcg 300
 gagaaatcac gacc 314

<210> 2290
 <211> 303
 <212> nucleic acid
 <213> Glycine max

 <400> 2290

 tcgcnacacg cgtacgtaag ctcggaattc ggctcgagat ttgaggccag gcaagcccca 60
 ctcaaccacc acacctctcc tegtacacgc tacccttttc tgctcttctt ctacctttca 120
 agtttttaaaa gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag 180
 ggacaccctg acaagctctg cgaccaaadc tccgatgctg tctcgcacgc ttgcctcgag 240
 caggaccacg acagcaaagt tgctgacgaa acatggcacc aaaaccaact tggatcatggt 300
 ctt 303

<210> 2291
 <211> 285
 <212> nucleic acid
 <213> Glycine max

 <400> 2291

 gtgcgatgca cgcgtacgta agctcgggaat tcggctcgag cttacaaca gcacaaagcg 60
 gggtactgtc tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa 120
 gttaaaatgg cccaagaaac ttctctattc acatctgaat cagtgaacga ggggcaccct 180
 gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240
 gacagcaagg ttgctgttnn aaacctgcac caagaccaac atggt 285

<210> 2292
 <211> 289
 <212> nucleic acid
 <213> Glycine max

 <400> 2292

 agtcgatgc acgcgtacgt aagctcggaa ttcggctcga ggcagactta acaacagcac 60
 aaagcggggtt actgtctgtt caagctacca tctctctctc tctttcttag tgctccttg 120
 ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180
 caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240
 gaccctgaca gcaaggttgc ctgtgaaacc tggaccaag accaacatg 289

<210> 2293
 <211> 343
 <212> nucleic acid
 <213> Glycine max

 <400> 2293

 acngcgtacg tnanctcgnn attcggtctg agggccaggc aagccccact caaccaccac 60
 acctctctc gttcacgcta cccctttctg ctcttcttct acctttcaag ttttaaaagt 120
 atacagatgg cagagacatt cctatttacc tcagagtgg tgaacgaggg acaccctgac 180
 aagctctgcg accaaatctc cgatgctgtc ctgcagcctt gcctcgagca ggacncagac 240
 agcaaagttg cctgcgaaac atgcacaaa accaattggt catggtcttc ggagaaatca 300

cgacagccan gttgatagag agatatgcgt gacnctgcag aca

343

<210> 2294

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2294

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accacacntc tctcgtttca cgctaccctt tctgtctctt cttctacctt tcaagtttta 120

aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180

ctgacancct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240

cagacagcaa agttgcctgc gaaacatgca ccanaaccaa cttgggtcat 289

<210> 2295

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2295

gtcgcangca cgcgtacgta nagctcggaa ttcggctcga gntgaggcca ggcaagcccc 60

actcaaccac cacacctctc ctcgttcacg ctaaccctt tctgtctctt ttctaccttt 120

caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180

agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctc 240

agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaacncac ttggtc 296

<210> 2296

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2296

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ccacacntct cctcgttcac gctaccctt tcttntctctt cttctacctt tncaagtttt 120

aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180

cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240

ccagacagca aagttgcctg cgaaacatgc accaaaacca acttgg 286

<210> 2297
<211> 318
<212> nucleic acid
<213> Glycine max

<400> 2297

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accaccacac ctctcctcgt tcacgctacc cttttctgct cttcttctac ctttcaagtt 120
ttaaagatat aaagatggca gngacattcc tatttacctc agagtcggtg aagagggaca 180
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240
cccagacagc aaagttgcct gcgaaacatg caccagaacc aacttgggtca tgggtcttcgg 300
agaaatcacg accaaggc 318

<210> 2298
<211> 290
<212> nucleic acid
<213> Glycine max

<400> 2298

ngnngcacgc gtacgtnagc tcggaattcg gctcgagttt gaggccaggc aagccccact 60
caaccaccac acgggctcct cgtnnacgct acccctttct ncctcttctt ctacctttca 120
agtttttaaaa gtataaaaat ggcagagaca ttctatttta cctcagagtc ggtgaacgag 180
ggacaccctg acaagctctg cgaccaaate tccgatgctg tcttcgacgc ttgcctcgag 240
caggacccag acagcaaagt tgctgcgaa acatgcacca aaaccaactt 290

<210> 2299
<211> 275
<212> nucleic acid
<213> Glycine max

<400> 2299

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aaagcggggtt actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg 120
ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180

caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240
gaccctgaca gcaaggttgc ctgtgaaacc tgcac 275

<210> 2300
<211> 308
<212> nucleic acid
<213> Glycine max
<400> 2300

ncgtcgang cagcgtacg taagctcgga attcggctcg agaggcaagc cccactcaac 60
cancacangg nnoctcgtn acgtacccc tntctnctc ttcttctacc tttcagngtn 120
ttaaagtat aaagatggca gagacattcc tatttaccag agtcggtgaa cgagggacac 180
cctgacaagc totgogacca aatctccgnt gctgtctcgc acgcttgccg cgagcaggac 240
ccagacagca aagttgcntg cgaaacatgc nccaaaacca acttggnccat ggtcttcgga 300
gaaatcag 308

<210> 2301
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2301

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aggccccant caaccaccac acctctanct cgttcacgct acccctttct gctcttcttc 120
taactttcaa gttttaaaag tataaagatg gcagagacat tcctatttac ctcagagtcg 180
gtgaacgagg gacaccctga caagctctgc gaccaaactc ccgatgctgt cctcgacgct 240
tgctcgagc aggaccaga cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg 300

<210> 2302
<211> 295
<212> nucleic acid
<213> Glycine max
<400> 2302

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ccaccacacc tctctcggtt cagctaccc ctttctgctc ttcttctacc tttcaagttt 120

taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacggcttgc ctcgagcagg 240
 acccagacag caaagttgcc tgcgaaacat gcancaaaac caacttggtc atggt 295

<210> 2303
 <211> 281
 <212> nucleic acid
 <213> Glycine max
 <400> 2303

atgcacgcgt acgtaagctc ggaattcggc tcgaggccat ttgggagtta ggttctgcac 60
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca nacgtgctt 120
 cttcngcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180
 gtcaccccgga caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccctgaaca 240
 ggaccctgac agcaagggttg cctgtgagac atgcaccaag a 281

<210> 2304
 <211> 297
 <212> nucleic acid
 <213> Glycine max
 <400> 2304

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 aactttccta ttcacatctg aatcagtga cgaggggcac cctgagaagt ctgtgaccag 180
 atctccgatg ctgtgctcga tgcattgntt gagcaggacc ctgacagcaa gggtgcctgt 240
 naaacctgca ncaagancaa catggtgatg gntttncgga gagatcacia acanngc 297

<210> 2305
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2305

gtgcacgcga cgcgtacgtn agctcggaat tcggctcgag cngccactaa cgtaatcagg 60
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ncgttaaaat ggcccaagaa actttcctat ncacatctga atcagtgaac gangggcacc 180
ctgacagctc tgtgaccaga tctccgatgc tgtgctcgnt gcntgcntgg agcaggaccc 240
tgacagcnag gntgcctgtg aaacctgcac caagacnnac atggtgangg ttttcggang 300
anatcacaac 310

<210> 2306
<211> 295
<212> nucleic acid
<213> Glycine max
<400> 2306

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagg gccaggcaag cccactcaa 60
ccaccacacc tctcctcggt cactctaccc ctttctgctc ttctcttacc tttcaagttt 120
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggacc 180
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct cgagcaggac 240
ccagacagca aagttgcctg cgaaacatgc accannacca acttggtcat ggtct 295

<210> 2307
<211> 158
<212> nucleic acid
<213> Glycine max
<400> 2307

cnagggacac cctgataagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct 60
cgaacaggac tcagacagca aggttgccctg cgaaacatgc accaagacca acttggtcat 120
ggtcttcgga gagacaccac caaggccaac gttgacta 158

<210> 2308
<211> 302
<212> nucleic acid
<213> Glycine max
<400> 2308

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gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120
agttaaaatg gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180

tgacaagctc tgtgancaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240
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<210> 2309
 <211> 295
 <212> nucleic acid
 <213> Glycine max
 <400> 2309

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 caccacacct ctctctgttc acgtacnnc tttctgctct tttctacct ttcaagtttt 120
 aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa cgagggacac 180
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
 ccagacagca aagttgcctg cgaaacatgc caccaaaaacc aacttggtca tggtc 295

<210> 2310
 <211> 271
 <212> nucleic acid
 <213> Glycine max
 <400> 2310

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 aagcgggtta ctgtctgttc aagctacat ctctctctct cttcttagt gcctccttgc 120
 cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc 180
 accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240
 accctgacag caaggttgcc tgtgaaacct g 271

<210> 2311
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2311

tnacangcac gcgtacgtaa gctcggatt cggctcgagg gccaggcaag cccactcaa 60
 ccaccacacc tctctcgtt cacgtaccc cttctgtctc ttctctacc tttcaagttt 120

taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240
 cccagacagc aaagttgcct gcgaaacatg caccanaacc aattgggtcat ggtttcggag 300
 aatcac 306

<210> 2312
 <211> 308
 <212> nucleic acid
 <213> Glycine max
 <400> 2312

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gcccactca 60
 accaccacac ctctcctcgt tcacgctacc cttttctgct cttctctac ctttcaagtt 120
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagnnggtga acgagggaca 180
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240
 cccagacagc aaagttgcct gcgaaacatg caccanaacc aacttggtca tggctcttcgg 300
 agaaatca 308

<210> 2313
 <211> 290
 <212> nucleic acid
 <213> Glycine max
 <400> 2313

acgtcgcang cagcgtacg taagctcgga attcggctcg ngcttgagca gacttaacaa 60
 cagcaciaaag cgggttactg tctgntcaag ctaccatctc tctctctctt tcttagtgcc 120
 tccttgccag aagttaaaat ggccaanna actttccnat tcacatctga atcagtgaac 180
 gaggggcacc ctgacaagct ctgtgaccag atctccgatg ctgtgctcga ngcatgcttg 240
 gagcaggacc ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa 290

<210> 2314
 <211> 294
 <212> nucleic acid
 <213> Glycine max
 <400> 2314

tcgcatgcac gcgtacgtaa gctcggaatt cncctcgagg caagccccac tcaaccacca 60
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctncagagtc ggtgaacgag ggacaccctg 180
acaagctctg cgaccaaate tccgatgctg tctctgacgc ttgcctcgag caggaccctg 240
acagcaaagt tgctgcgaa acatgcacca aaaccaactt ggtcatgggc ttcg 294

<210> 2315
<211> 297
<212> nucleic acid
<213> Glycine max
<400> 2315

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ccccactcaa ccaccacacc tctcctcggt cagcgtaccc ctttctgctc ttcttctacc 120
tttnaagttt taaaagtata aagatggcag agacattcct atttacctca gagtccgtga 180
acgaggggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240
tcgagcagga ccagacagc aaagttgcct gcgaaacatg caccaaaacc aantggt 297

<210> 2316
<211> 233
<212> nucleic acid
<213> Glycine max
<400> 2316

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acgagggggca ccctgacaag ctctgtgncc agatctccga tgctgtgctc gatgcatgcn 180
tgagcagga ccctgacanc aaggttgctt gtgaaacctg caccaagacc anc 233

<210> 2317
<211> 288
<212> nucleic acid
<213> Glycine max
<400> 2317

tcgtgcacg cgtacgtaag ctcggaattc ggctcgaggc caggcaagcc cactcaacc 60

accacacctc tctctgttca cgctaccctt tttctgtctt cttctacctt tcaagtttta 120
aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240
cagacagcaa agttgcctgc gaaacatgca ccaaaaccaa ttggtcat 288

<210> 2318
<211> 304
<212> nucleic acid
<213> Glycine max
<400> 2318

nnctgtcgan gcaecgttac gtaagctcgg aattcggctc ganactcgag ccgattcggc 60
tcgagggccc cactcaacca ccacacctct anctcgttca cgctaccacc tttctgtctt 120
tcttctaaact ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag 180
agtcggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg 240
acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca 300
attg 304

<210> 2319
<211> 305
<212> nucleic acid
<213> Glycine max
<400> 2319

tcgcangcac gcgtacgtna gctcggcnatt cggctcgagn ntttgaggcc aggcaagccc 60
cactcaacca ccacacctct cctcgttcac gctaccctt tctgtcttct ttctaccttt 120
caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180
agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctg 240
agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaan tgggtcatggt 300
cttcg 305

<210> 2320
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 2320

cgcatgcacg cgtacgtaag ctcggaattc ggctcgaggg ccaggcaagc ccactcaac 60
caccacacct ctctcgttc acgtacccc tttctgctct tttctacct ttcaagtttt 120
aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
ccagacagca aagttgcctg cgaaacatgc accaaaacca attggtcatg gtcttcgga 299

<210> 2321

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 2321

tcgcatgcac gcgtacgtna gctcggaatt cngctcgagc ngctcgagca cacctctcct 60
cgttcacgct acccctttct gctcttcttc tacctttcaa gttntaaaag tataaagatg 120
gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacacctgac aagnctgcga 180
ccaaactccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc 240
tgcgaaacat gcacaaaaac caattgggtca tggctctcgg agaaatcacg accaggccaa 300
gttgactacg agaaga 316

<210> 2322

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2322

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tctgcacgct ctgcttcag cgagtgttct ttcttcgttt caacacctta atttgcacac 120
gctgcttctt cagcttgaga aatggcacia gaaacctttc tattcacatc tgaatctgta 180
aacgagggtc accccgacaa gctgtgcgac cagatctctg atgagtgtc gatgcgtgcc 240
ttgaacagga cctgacagc aaggttgctt gtgagacatg caccaaga 288

<210> 2323

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2323

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 acanctctcc tcgttcacgc tacccttnc tgctcttctt ctacctttca agtttttaaaa 120
 gtataaagat ggcagagaca ttinctattta cctcagagtc ggtgaacgag ggacaccctg 180
 acaagctctg cgaccaaate tccgatgntg tccctgaatg naaatcgagc aggaccaga 240
 nagcaaagtt gntgcgana catgcaccaa aaccaacttg gtcatggtct tcggagaaa 299

<210> 2324

<211> 254

<212> nucleic acid

<213> Glycine max

<400> 2324

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 tgcctccttg ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt 120
 gaacgagggg caccgcacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc 180
 ttggagcagg accctgacag caaggttgcc tgtgaaacct ngcaccaaga ccnacatggt 240
 gatggttttc ggag 254

<210> 2325

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2325

cacgcgtacg taagctcgga attcggctcg annttctcaa cattgtgaag gaaaactttg 60
 atttcaggcc tggatgata tccatccaac cttgatctca agaggggtgg aaataacagg 120
 tttttgaaga ctgctgcta tggacacttt ggaagagaag accctgacnt tcacatgggg 180
 aagttggtcc naccctctcn agttgggnaa agccnaacca tttcatc 227

<210> 2326

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2326

gtcatgnagg acgnangacg tgnagctcag nattcggntc gagcggttcac gctaaccct 60
ttctggctct ncttcnanc tccaagtttt aaaagtatan agatgngcag aganattcct 120
atttacctac agagtcggtt aacgagggac accctgacaa gctctgacgac caaatcccga 180
tgctntcttc gagcgttgcc tcgagcngga cccagacagc aaagttgcct gcgaaacatg 240
caccaaaacc aacttggtca tgggtcttcg aganntcang accaaggcca acgt 294

<210> 2327

<211> 281

<212> nucleic acid

<213> Glycine max

<400> 2327

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anctaccatc tctctctctc tntcttagtg cctcccgtn cagnngtnan aatggcccaa 120
naaactttcc tattnacatc tgaatcagtg aacgangggc anccganaan ctccgtgacc 180
agatntccga tgctgtgntc gatgcatgct tggagcagga cctgacagc aaggttgct 240
gtganacctg caccaagacc ancatggtgn tggtttncgg a 281

<210> 2328

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2328

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gtctgttcaa gctaccatcc tctctctctc nttcttagtg cctcctnnn anaagttaa 120
atggccann anactttcct atncacatct gaatcantga acgaggggca cnctgacaag 180
ntctgtgncc agatctccgg tgctgtgctc gatgcatgct tggagcagga cctgacagc 240
aagntgcct ggaaacctgc acnaagacca acatggtgat ngntttcgga gann 294

<210> 2329

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2329

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tcaaccacca cacctctcct cggtcacgct acccctttct gctacttctt ctacctttca 120
agtttttaaaa gtataaagat ggcagagaca ttcctattta cctcagagtc ggtgaacgag 180
ggacaccctg acangctctg cgaccaaatt tccgatgctg tcctcgacgc ttgcctcgag 240
caggacccag acagcaaagt tgctgcgaa acatgcacca aaac 284

<210> 2330

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2330

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aaccaccaca cctctcctcg ttcacgctac cctttctgc tcttcttcta cctttcaagt 120
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggga 180
caccctgaca agotctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240
gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaa cc 282

<210> 2331

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 2331

angtcgcang caocggtacg taagctcgga attcggtcgc nctcgagcc gaatnggctc 60
gagcacaaaag cgggntactg ncngntacaa gctacnatct ctncctctc tntcttagtg 120
nctacctgcc agangtnaaa atggcnaag aaactntnct attnnnatct gnttcagtga 180
acgaggggca cnetgacaan ntctttgacg agatctccga tgctgtgntc gatgcatgct 240
tggagcagga ccctgacagc aaggttgctt gtnaaacctg cncnaaananc aacatggtga 300
tggtttcgg 309

<210> 2332

<211> 262

<212> nucleic acid

GeneBank

<213> Glycine max

<400> 2332

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gtntttttgt anntttcaag ntttaaaagt ataaagatgg cagagacatt cctatttacc 120
tcagagtcgg tgaacgaggg acaccctgac aagctntgcg accaaatctc cgatgctgtc 180
ctcgacgctt gctcagagca ggaccagac agcaaagttg cnngcgaaac atggcaccaa 240
aaccaattgg tcatggtntt cg 262

<210> 2333

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2333

tngtctcatg cagcgtacg taagctcgga attcggtctg anctcgancc gnttcggcta 60
ctagaagccc cactcaacca ncacacctcn cctcgttca ngctaccctt ttctgctctt 120
cttctacctt caagttttta aagtataaag atggcagaga catttctatt tacctcagag 180
tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac 240
gcttgccctg agcaggaccc agacagcaa gttgcctgcg aaacatgcac caaaac 296

<210> 2334

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2334

tngtcgcang cagcgtacg taagctcgga attcggtctg nnctcgagcc gatttcggctc 60
gagcaacagc acaaagcggg ttactgtngg ttcaagctac catctctact ctctctntct 120
tagagcctcc ttgccagaag ttaaaatggc ccaagaaact ttctatttca catctgaatc 180
agtganogag gggcaccttg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 240
atgcttggag caggaccctg acagcaaggt tgcctgtgaa acctgcacca a 291

<210> 2335

<211> 271

<212> nucleic acid

<213> Glycine max

<400> 2335

cgtcgttgca cgcgtacgta agctcggaat tcggctcgcn agctccactc aaccaccaca 60
 cntctcctcg ttcacgctac ccctttctgc tctttctcta cctttcaagt tttaaaagta 120
 taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca 180
 agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
 gcaaagttgc ctgcgaaaca tgcacaaaaa c 271

<210> 2336

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2336

gtgcgancga cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gcccactca 60
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 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaattct cgatgctgtc ctgacgctt gcctcgagca 240
 ggaccagac agcaaagttg cctgcgaaac atgcacaaaa acca 284

<210> 2337

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2337

ncgtcgancg cacncgtacg taagctcgga attcggctcg nctcgagcc gattcggctc 60
 gagcaagccc cactgcaacc accacaacnt cttcctcgtt cagactacc cctttctgct 120
 cttcactcta cctttcaagt tttaaaagta taaagatggc agagacattc ctatttacct 180
 cagagtcggt gaacgagggg caccctgaca agctctgcga ccaaattctc gatgctgtcc 240
 tcgacgcttg cctcgagcag gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa 300
 cca 303

<210> 2338

<211> 183
 <212> nucleic acid
 <213> Glycine max
 <400> 2338
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 cctcagccat gtccttgcaa ccaaactcgg tgctcgctc accgagggtta ggaaaaatgg 120
 tacctgtgct tggctgaggc cagatggcaa gacacaagta actgttgagt actacaatga 180
 caa 183

<210> 2339
 <211> 322
 <212> nucleic acid
 <213> Glycine max
 <400> 2339
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 gggcagcgct tgatttgagg ccaggcaagc ccactcaac caccacacct ctctcgttc 120
 acgctacncc tttctgntct tcttctacct ttcaagtttt aaaagtataa agatggcaga 180
 gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 240
 aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcccg 300
 cgaaacatgc accaaaacca ac 322

<210> 2340
 <211> 259
 <212> nucleic acid
 <213> Glycine max
 <400> 2340
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 ngctgtgtna agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaa 120
 atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180
 ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240
 aaggttgctt gtgaaacct 259

<210> 2341

<211> 309
 <212> nucleic acid
 <213> Glycine max

 <400> 2341

 ncgcatgcac gcgtacgtaa gctcgggnatt cngctcggnnc tcgagccgat tcggctcgag 60
 aaacgagcca tttggnagtn aggtncctgc acgntctgct taccgcgnag tgttctttct 120
 tcacttcaac accttaattt ncacacgctg cttcttcngc ttgaganatg gcacaagaaa 180
 cntttctatt cacatctgaa tctgtaaacg anggtcaccc cgacaagctg tgcgaccaga 240
 tctctgatgc agtgctcgat gcgtgccttg aacaggaccc tgacagcaag gttncctgtn 300
 agacatgen 309

<210> 2342
 <211> 277
 <212> nucleic acid
 <213> Glycine max

 <400> 2342

 cgtgcagcgt acgtaagctc ggaattcggc tcgagcagac ttaacaacag cacnaagcgg 60
 gttantgtct gttcaagcta ccattctctc ctctntttct tagtgctcc ttgccagang 120
 ttaaaatggc ccaanaaact ttctntttca catctgaatc agtgaacgag gggcaccctg 180
 acaagntctg tgaccagatc tccgntgctg tgctcgatgc atgcttgag caggaccctg 240
 acancaaggt tgctgtgan acctgcacca agaccaa 277

<210> 2343
 <211> 287
 <212> nucleic acid
 <213> Glycine max

 <400> 2343

 gtgcangca cgcgtacgtn agctcggnat tcggctcgag gnggccaggc aagccccact 60
 caaccaccac acctctctc gttcacgcta cccctttctg ctcttcttct acctttcaag 120
 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ttcgacgctt gcctcgagca 240
 ggaccagac agcaaagttg cctgcgaaac atgcacaaa nccaact 287

<400>	2344
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[illegible]

<400> 2345

[illegible]

<400> 2346

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 cgtgcctgan caggaccctg acagcaaggt tgccctgtgag acatgcacca agaccaantg 300
 gtcattgg 307

<210> 2347
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2347

gtcnctgcac gcgtacgtaa gctcgggaatc ggctcgannc tcgagccgaa tcggctcgag 60
 aacagcacia agcggggttac tgtcacttca agctaccatc tatctctnln ctttcttagt 120
 gntccttgg ccanaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt 180
 gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatggcat 240
 gcttggagca ggaccctgac agcaagggtt nctgtgnaaa cntggtagca agaccaacat 300
 ggtgatgggt ttcggn 316

<210> 2348
 <211> 281
 <212> nucleic acid
 <213> Glycine max
 <400> 2348

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 cactctctct gctcttcttc tacctttcaa gtttttaaag tattaagatg gcagagacat 120
 tcctatttac ctgagagtca gtgaacgagg gacaccctga cagctctgag accaaatctc 180
 cgatgctgtc ctgacgctt gccttgaaca ggaccagac ancaagggtg cctgcgaaac 240
 atggcaccaa gaccaattgg tcattggtctt cggagagatc a 281

<210> 2349
 <211> 295
 <212> nucleic acid
 <213> Glycine max
 <400> 2349

aaagtcgan gcacgcgtac gtaagctcgg aattcggctc gagcactcaa ccaccacacc 60

tctcctcggt cagcgtaccc ntttctgnct ncttcttcta cctttcaagt tttaaaagta 120
 taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca 180
 agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
 gcaaagttgc ctggcgaaac atgcaccaa accaacttgg tccatggtct tcgga 295

<210> 2350
 <211> 491
 <212> nucleic acid
 <213> Glycine max
 <400> 2350

tttacttnnc ccgnccggta caagtcataa attcccgggn cgacccaagc ntccnnccac 60
 gcgctcggtac ggctgcgaga agacgacaga agggggcagc gcttgatttg aggccaggca 120
 agccccactc aaccaccaca cctctctctg ttcacgtac ccctttctgc tcttcttcta 180
 cctttcaagt tttaaaagta taaagatggc agnacattc ctatttacct canagtcggt 240
 gaacgagggg cncctgaca agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg 300
 ccnecatcag gaccagaca gcaaanttgc ctgcaaaaaca tgcacaaaa accaacttgg 360
 tcatggtctt cggagaaatc angaccaa anccaacgtng actacngaag aanaatagt 420
 cgttnacacc tngcaggaac atcngcttc gtctcaaaat gatttgngga ctggattcnc 480
 naacaactgg g 491

<210> 2351
 <211> 276
 <212> nucleic acid
 <213> Glycine max
 <400> 2351

gtcgnangan gcacgcgtac gtnagctcgg aattcggctc gagnacgac aaagcgggtt 60
 acngtctgtt caagctacca nctctctctc tctttcttag tgcctccttg ccagangtta 120
 aaatggccca agaaacttnc ctannacat ctgaatcagt gaacgagggg caccctgaca 180
 agctcngtga ccagatcncc gatnctgtgc ncgatgcatg cttggagcag gaccctgaca 240
 gcaaggtngc ctgtgaaanc tgcaccaaga ncaact 276

<210> 2352

<211> 315
 <212> nucleic acid
 <213> Glycine max

 <400> 2352

 anggnngngc acgcgtacgt aagctcggaa ttcggtcga gnggenagcc ccaactcaacc 60
 accacacctc tctcgttca ngctaccctt ttctgtctt cttctacctt tcaagtttta 120
 aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctc gagcaggacc 240
 cagagagcaa ggttgcntgc gaaacangca ccnaaaanca anttggtgat ggtccttcgg 300
 ngattcacga ccang 315

<210> 2353
 <211> 287
 <212> nucleic acid
 <213> Glycine max

 <400> 2353

 gconangcac gcgtacgtaa gctcggaatt cggctcgagt gatttgangc taggcaagcc 60
 ccantcaacc accacacctc tctcgttca cgctaccctt ttctgtctt cttctacatt 120
 tonagtttta aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac 180
 gtnggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctc 240
 gagcaggacc cagacagcaa agttgcctgc gaaacatgca ccaaaac 287

<210> 2354
 <211> 277
 <212> nucleic acid
 <213> Glycine max

 <400> 2354

 agtcgcangc acgcgtacgt aagctcgga ttctngctcga gggcaagccn cactcaacca 60
 cnacacntct cctcgttnca cgctaccctt ttgtctctt tcttctacct ttcaagtttt 120
 aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
 ccagacagca aagttgcctg cgatacatg accaaaa 277

<210> 2355
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 2355

 cnntcaacnn gnanagtcgc angcacgcgt acgtaagctc ggaattcggc tcgaggcaga 60
 ettaacaaca gcacaaagcg gggtactgtc tgttcaagct accatctctc tctctctttc 120
 ttagtgcttc cttgccagaa gttaaaatgg cccaagaaac tttcctattc acatctgaat 180
 cagtgaacga ggggcaccct gacaagctct gtgaccagat ctccgatgct gtgctcgatg 240
 catgcttgga gcaggaccct gacagcaagg ttgcctgtga aacctggaac caaggnocan 300
 tttgnn 306

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<210> 2356
 <211> 285
 <212> nucleic acid
 <213> Glycine max

 <400> 2356

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 caaccaccac acctctcttc gttcacgcta cccctttctg ctctttctctt acctttcaag 120
 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgncgctt gcctcgagca 240
 ggaccagac agcaaagttg cctgcgaaac atgcacaaaa ancaa 285

<210> 2357
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2357

 nagtcgcang cacgcgtacg taagctcgga attcggctcg agttgaggcc aggcaagccc 60
 cactcaacca ccacacctct cctcggtcac gctaccctt tctgctcttc ttctaccttt 120
 caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180
 agggacaccc tgacaagctc tgcgacaaa tctccgatgc tgtcctcgac gcttgccctg 240

agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaacaact tggtcatg 298

<210> 2358
 <211> 288
 <212> nucleic acid
 <213> Glycine max
 <400> 2358

gtngcangca cgcgtacgta agctcggaat tcggctcgag nttgaggcca ggcaagcccc 60
 antcaaccac canacacnct cctcgtnnat ggctaccctt ttctnnotct tcttgetacc 120
 titncaagtt tnaaaagtat aaagatggca ganacattcc tatttacctc agagtcggtn 180
 aacgagggac accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc 240
 ctcgagcagg acccagacag caaagttgcn tgcgaaacat gcaccaan 288

<210> 2359
 <211> 335
 <212> nucleic acid
 <213> Glycine max
 <400> 2359

tcgcatgcac gcgtacgtaa gctcggaatt cngctcgagg caagccccac tcaaccacca 60
 cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
 tatanagatg gcagagacat tcctatttac ctcagagtcg gtgaacgngg ngacaccctg 180
 acaagctctg cgaccaaata tccgatgctg tctcgcgcg ttgcctcgag caggacccag 240
 acagcaaagt tgcttgcgaa acatgcacca aaaccaactt ggtcatgggc ttccgagaaa 300
 tcacgaccaa gccaaattga ctagaagaag atatg 335

<210> 2360
 <211> 505
 <212> nucleic acid
 <213> Glycine max
 <400> 2360

ttcnaacttt caccgcccag gtaacngtan aagaattccc ggnccgaccc acnggtcanc 60
 ccangcgtcc gccacgcgt ccgtacggct gcnanaagac gacagaangg gcaccgcttg 120
 agcagactta acaacatnac aaaccgggtt actgtctgtt canggctacc atctctctct 180

ctctttcttta ggtgnctcct tgccagnang tnnnaatgng gcnaagnaac ttncctatct 240
 acatctgant cagtgaacga tgggnaccnt gacaagctct gtgaccagat cnccgatgcn 300
 gtgctcgatg cangcttgga gcaggaccct gacagcnncg ttgcctgtga aacctgcacc 360
 aataccanca tggatgatggg ttttcggaga gatnccaanc gangccnaan nttgnaacta 420
 ttaggaagat tngnggcgnt gacncatnca ggaananttn ctttgncgcc nngntnatgg 480
 ntgcctctgn nggcnganaa ncttc 505

<210> 2361
 <211> 283
 <212> nucleic acid
 <213> Glycine max

<400> 2361
 acgtcgang cagcggtacg taagctcgga attcggtctg aggcagactt aacaacagca 60
 canagcgggt tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt 120
 gccagaagtt aaaatggccc aagaancctt cctattcaca tctgaatcag tgaacgaggg 180
 gcacctgac aagctctgtg accagatctc cgatgctgtg tcgatgcatg cttggagcag 240
 gacctgaca gcaaggttgc ctgtgaaacc tgcaccaaga cca 283

<210> 2362
 <211> 495
 <212> nucleic acid
 <213> Glycine max

<400> 2362
 tttacnagct cttacgcggn caantaccgt gtntaggnat tcccgggctc gacncacgog 60
 tcagtacggc tgcgagaagn gaacnnaggg ggcaacacnt ggnttgangn catgcaagcc 120
 ccaactcaacc accacacctc nctcgttca cgctaccctt ttctgtctt cttctanctt 180
 tcatntttta anncgatan agatggcana gacattccta tttacntcag ngtcagtga 240
 cganggacac nctgacaagc tctgcgagca aatctccagn tgctgtcctc gacgcttgcc 300
 tcgancagga ccatacagc aaacttgctt gccaanatg ctccaaaacc ancttggtcn 360
 tngtcttcgg anaaatcn cn accaaggcca acnttgactn cnanangata ntgcgttaca 420
 ctgcnggaac atnggcttct tctcatatga tntgggactg gattccnact cctgcaaagt 480

495

<400>	2363
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[illegible]

<400> 2364

[illegible]

<400>	2365
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832

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<210> 2366
 <211> 316
 <212> nucleic acid
 <213> Glycine max

<400> 2366

naaaanaang gaaantcgca tgcacgcgta cgtnagctcg gaattcggct cgagggccag 60
 gcaagcccca ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt 120
 ctacctttca agtttttaaaa gtataaagat ggcagagaca ttctatttta cctcagagtc 180
 ggtgaacgag ggacaccctg acaagctctg cgaccaaata tccgatgctg tctcgcagcg 240
 ttgcctcgag caggacccag acagcaaagt tgcctgcgaa actgcaccan aaccaattgg 300
 tcatgggtctt cggaga 316

<210> 2367
 <211> 289
 <212> nucleic acid
 <213> Glycine max

<400> 2367

gtcgcngcac gcgtacgtga gctcggaatt cggctcgagg gccaggcaag ccccaactcaa 60
 ccaccacacc tctctctggt cacgctaccc ctttctgctc ttcttctacc tttacaagtt 120
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcgggtg aacgagggac 180
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240
 accagacag caaagttgcc tgcgaaactg caccancacc aattgggtca 289

<210> 2368
 <211> 302
 <212> nucleic acid
 <213> Glycine max

<400> 2368

gtcgcangca cgcgtacgtn agctcggaat tcggctcgag ggccaggcaa gcccactca 60
 accaccacac ctctctctgt tcacgctacc ctttctgctc cttctctac ctttcaagtt 120
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggta acgagggaca 180

ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240
 cccagacagc aaagttgcct gcgaaacatg caccataacc aacttggtca tgggtcttcgg 300
 ag 302

<210> 2369
 <211> 288
 <212> nucleic acid
 <213> Glycine max
 <400> 2369

cgcatgcacg cgtacgtnag ctcggaattc ggctcgagtt tgaggccagg caagcnccac 60
 tcaaccacca cacctctcct cgttcacgct acccctttct gctacttact tctacctttc 120
 aagtttttaa agtataaaga tggcagagac attcctattt acctannagc cgggtgaacga 180
 gggacaccct gacaagctct gcgaccaaatt ctccgatgct gtcctcgacg cttgctctga 240
 gcaggaccca gacagcaaag ttgcctgcga aacatgcacc aaaacnaa 288

<210> 2370
 <211> 292
 <212> nucleic acid
 <213> Glycine max
 <400> 2370

cgcangcacg cgtacgtaag ctcggaattc ngctcgagca agccccactc aaccaccaca 60
 cctctcctcg ttcacgctac ccctttctgc tcttcttcta cctttcaagt tttaaaagta 120
 taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggga caccctgaca 180
 agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240
 gcaaagttgc ctgcgaaact gcacaaaaac caatgggtca ngntctnaga aa 292

<210> 2371
 <211> 288
 <212> nucleic acid
 <213> Glycine max
 <400> 2371

gtcgngcac gcgtacgtaa gctcgggnatt cngctcgagg caagccccac tcaaccacca 60
 cacctctacc tacgttcacg ctaccccttt ctgctcttct tctaccttta caagttttaa 120

aagtataaag atggcagaga cattcctatt tacctacaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240
 cagacagcaa agttgcctgc gaaacatgca ccaaaaccaa cttggtca 288

<210> 2372
 <211> 299
 <212> nucleic acid
 <213> Glycine max

<400> 2372

acgtcgctgc acgcgtacgt aagctcggaa ttcggtcga ggtgatttgg agtttggagc 60
 gactgaacta ntcattaatt tgcacttcgc tgtttcagct tcatcaccct tcttttgcac 120
 catttatatc tcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180
 ccanggtcac cccgacaagc tgtgcgacca gatctctgat gcagtaacttg atgcntgcct 240
 gnncaggacc ctggacagcn aggtancctg tnagacatgc accaagacca acatggtca 299

<210> 2373
 <211> 283
 <212> nucleic acid
 <213> Glycine max

<400> 2373

cgcatgcacg cgtacgtnag ctcggaattc ggctcgaggg caagccccac tcaaccacca 60
 cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
 tataaagatg gcagagacat tcctatttac ctacagantcg gtgancgagg gncaccctga 180
 caagctctgc gaccaaattct ccgatgctgt cctcgacgct ngcctcganc aggaccana 240
 cagcnaagtt gcctncgana catgnaccaa aaccaacttg gtc 283

<210> 2374
 <211> 283
 <212> nucleic acid
 <213> Glycine max

<400> 2374

gannagtgc angcacgcgt acgtaagctc ggaattcggc tcgaggnaca gcacaaagcg 60
 ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgcctc cttgccagaa 120

gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180
gacaagctct gtgaccagat ctccgatgct ntgctcgatg catgcttgga gcaggaccct 240
gacagcaagg ttgcctgtga aaactgcaac caagaccaan nat 283

<210> 2375
<211> 302
<212> nucleic acid
<213> Glycine max
<400> 2375

nanncgchang cacgcgtacg taagctcgga attcggctcg agctcgagcc gtggagtttg 60
gagcgactga actaatcatt aatttgcact ncgctgtttc agcttcatca cccttctttt 120
gcatcattta tatctcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180
taaachnaggg tcaccccgac aagctgtncn accagatctc tgatgcagta cttgatgcgn 240
gcctgancag gaccctgaca gcaagggttg ctgtgagaca tgcaccaagn cccaacatgg 300
tc 302

<210> 2376
<211> 270
<212> nucleic acid
<213> Glycine max
<400> 2376

gcannngtnac gtaagctncg gaattcggct cgaggactta acaacagcac aaagcggggtt 60
actgtgctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgaa 180
caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga 240
cagcaagggtt gcctgtgana cctgcaccaa 270

<210> 2377
<211> 312
<212> nucleic acid
<213> Glycine max
<400> 2377

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gagaccaagc cccactcaac 60

cancacacca ntctactctg ctcttcttct acctttcaag tttctaaagt atnaagatgg 120
cagagacatt cctattnacc tcagagtcag tgaacnaggg acacctgat caagctctgc 180
gagccanac tccgatgctg tctctgacgg cttgccttga acaggaccca gacagcaagg 240
ttgcctgcga aacatgcacc aagaccaact tggtcatggg cttcggagag atcaccncca 300
aggccaacgt tg 312

<210> 2378
<211> 328
<212> nucleic acid
<213> Glycine max
<400> 2378

agtcgcatgc acncgtacgt aagctcggaa tttcggctcg aggcattgtg gcaagtggac 60
tagccagaag gtgcancncn gcaagtgtct tatgccattg gtgtgcncga ncctttgtnc 120
tgtatttgtt gacacctatg gcaccgggaa gatccatgat aaggagattc tcaacattgt 180
gaaggagaac ttgatttcan ncccggatg atctcccatc aaccttgatn tcaagagggg 240
tggaataac aggttcttga agatgctgca tatggacatt cggcagagag ncgnattcac 300
aggggatggg cnangcccc ccaatggg 328

<210> 2379
<211> 258
<212> nucleic acid
<213> Glycine max
<400> 2379

cagcacaang cnggttacng ncnngtcaag cnaccatctc tctctctctt tnttagtggc 60
ctccttgcca gaagttaaaa tngcccaaga aactttcnta ttcacatctn aanccagtna 120
acgaggggca cctgacaag ctctgtganc agatcnccga tgcngtgctc gntgcatnnt 180
tggagcanga cctgacagc aaggttgcct gtgaaacctg naccaanacc aacatggnga 240
ngggttcggg gagatcac 258

<210> 2380
<211> 267
<212> nucleic acid
<213> Glycine max

<400> 2380

gtcgcanang angcgnacgt ncagctcgga attcggtctg aggtccggta tgatctccat 60

caaccttnga tctcaagagg ggtgggaata acagggttctt gaagactgct gcatatggac 120

acttcggcag agaggaccct gacttcacat gggaagtggc caagcccctc aagtgggaga 180

nggcctaagg ccattcattc cacngcaatg tgctgggagt ttttnagcgt tgcccttata 240

atgncatatta tccataactt tccacgt 267

<210> 2381

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2381

agtcnnatna acgcgtacgt aagctcggaa ttcggtctga gtacggctgc nagaagacga 60

cagaaggggg cagcgcttga tttgaggcca ggcaagcccc actcaaccac cacacctctc 120

ctcgttcacg ctaccoccttt ctgctcttct tctacctttc aagtttttaa agtataaaga 180

tggcagagac attcctatatt acctcagagt cgggtgaacga gggacaccct gacaagctct 240

gcgaccaaatt ctccgatgct gtcctcgacg cttgcctcga gcaggaccca gacagcaaag 300

ttgcctgcga a 311

<210> 2382

<211> 235

<212> nucleic acid

<213> Glycine max

<400> 2382

tttgaggcca ggcaagcccc actcaaccac cacacctctc ctcgttcacg ctaccoccttt 60

actgctctnc ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt 120

tacctcagag tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc 180

gtcctcgac gcttgctcgc agcaggaccc agacagcaaa gttgcctgcg aaaca 235

<210> 2383

<211> 168

<212> nucleic acid

<213> Glycine max

<400> 2383

ncngcncgng tacnggtacg cnagctcgga attcggtcg aggtgctggt gaccagggtc 60
acatgtttgg ctatgccact gatgaaaccc ctgaattgat gccattgagc catgttcttg 120
caacaaaact cgggtgctcgt ctcaccgagg ttcgnaagaa cggtaacct 168

<210> 2384

<211> 156

<212> nucleic acid

<213> Glycine max

<400> 2384

tagtgcctcc ttgccagaag ttaaaatggc ccaagaaact ttctattca catctgaatc 60
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttgag caggaccctg acagcaaggt tgctg 156

<210> 2385

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 2385

ggccaggcaa gcccactca accaccacac ctctcctcgt tcacgctacn cttttctgct 60
cttctctac ntttcaagtt ttaaaagtat aaagatggca gagacattnc ctatttacct 120
canagtcggt gaacgaggga caccctgaca agctctgcga ccaaattctcc gatgctgtcc 180
tcgacgcttg cctcgagcag gactcagaca gcaaagttgc ctgcgaaaca tggcaccaan 240
accaattggt catggtcttc ggagaatcac gnccaagg 278

<210> 2386

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 2386

gtcgcangca cgcgtacgtn agctcggnat tcggtcgcag ggccaggcaa gcccactca 60
accaccacac ctctcctcgt tcacgctacc cttttctgct cttctctac ctttcaagtt 120
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgaggag 180

accctgacaa gctctgacgac caaatctccg atgctgtcct tcgacgcttg cctcgagcag 240
gacccagaca gcaaagttgc ctgcgaaaca tgcaccaa 278

<210> 2387
<211> 309
<212> nucleic acid
<213> Glycine max

<400> 2387

nngtcgcang cacgcgtacg taagctcgga attcggtcgc agtacggctg cnagaagacg 60
acagaagggg gcagcgttg atttgaggcc aggcaagccc cactcaacca ccacacctct 120
cctgcgttca gctacccctt tctgctcttc ttctaccttt caagttttta aagtataaag 180
atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc tgacaagctc 240
tgcgacaaa tctccgatgc tgtcctcgac gcttgctcgc agcaggaccc agacagcaaa 300
gttgctgc 309

<210> 2388
<211> 219
<212> nucleic acid
<213> Glycine max

<400> 2388

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tctctttctt agtgctcctt tgccagaagt taaaatggcc caagaaactt tcctattcac 120
atctgaatca gtgaacgagg ggcacctga caagctctgt gaccagatct ccgatgctgt 180
gctcgatgca tgcttggagc aggacctga cagcaaggt 219

<210> 2389
<211> 314
<212> nucleic acid
<213> Glycine max

<400> 2389

gngtacgtna gctcggaatt cggctccgag ngatttgagg ccaggcaagc cccactcaac 60
caccacacnt ctctcgttc acgctacccc tttctgctct tcttctacct ttcaagtttt 120
aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180

cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240
ccagacagcn aagttgcctg cgaaactgca ccaaaaccaa tnggtcaggt cttcggggaat 300
cngacaagcc acgt 314

<210> 2390
<211> 287
<212> nucleic acid
<213> Glycine max

<400> 2390

ggaannntcg cangcacgcg tacgttagct cggaattcgg ctcgagnact taacaacagc 60
acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt agtgctctct 120
tgccagaagt taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg 180
ggcacctgac aagctctgtg accagatctc cgatgctgtg ctgatgcat gttggagcag 240
gacctgacag caaggttgcc tgtgaaacct gcaccaagac caanatg 287

<210> 2391
<211> 281
<212> nucleic acid
<213> Glycine max

<400> 2391

annantcgca ngcacgcgta cgtnagctcg gaattcngct cgagggccag gcnagcccca 60
ctcaaccacc acacnactcc tcnttcncgc tacccttttt ctgcnctct tctacctttc 120
aagtttttaa agtatanaga tggcagagac attcctatct acctcagagt cggatgaacga 180
gggacaccct gacaagctct ggcaccaaatt ctccgatgct gtgcttcgac gcttgctctg 240
agcaggacnc agacagcaaa gttgcttcg aaacatgcac c 281

<210> 2392
<211> 275
<212> nucleic acid
<213> Glycine max

<400> 2392

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc ccaactcaacc 60
accacacctc tcctcggttca cgctaccctt ttctgctctt cttctacctt tcaagtttta 120

aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gaggacaccc 180
 tgacaagctc tgcgaccaa tctccgatgc tgtctcgcac gcttgcntcg agcaggaccc 240
 agacagcaaa gttgcctgcg aaacatgcag caaaa 275

<210> 2393
 <211> 303
 <212> nucleic acid
 <213> Glycine max

<400> 2393

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagt acggctgcga gaagaccaca 60
 gaagggggca gcgcttgaga ccaagcccca ctcaaccacc acaccactct ctctgctctt 120
 cttctacctt tcaagttttt aaagtattaa gatggcagag acattcctat ttacctcaga 180
 gtcagtgaac gaggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga 240
 cgcttgctt gaacaggacc cagacagcaa ggttgctgga aacatgnana agnccatntg 300
 ggt 303

<210> 2394
 <211> 189
 <212> nucleic acid
 <213> Glycine max

<400> 2394

cgtaagctcg gaattcggct cgaggtttca acaccttaat ttgcacacgc tgctttctca 60
 gcttgagaaa tggcacaaga nacctttcta ttcacatctg aatctgtaaa cgagggtcac 120
 cccgacaagc tgtgcgacca gatctctgat gcagtgcctg atgcgtgcct tgaacaggac 180
 cctgacagc 189

<210> 2395
 <211> 183
 <212> nucleic acid
 <213> Glycine max

<400> 2395

gctgcacgcg tacgtaagct cggaattcgg ctgcagctca agtttttgaa gtatagagat 60
 ggcagagaca ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg 120

tgaccaaadc tctgatgctg tctctgacgc ttgcctcgaa caggacccag acagcaaggt 180
tgc 183

<210> 2396
<211> 292
<212> nucleic acid
<213> Glycine max
<400> 2396

gtcgcangca cgcgtacgtn nagctcggna ttccgctcgn cctctgagcc gaattcgggt 60
cgagcaagcc ccaactcaacc accacaccnc tctctgttca cgcacncccc tttctgtctt 120
tcttccacct ttcaagtttt aaaagtataa agatggcaga gacattcctt ttacctcaga 180
gtcgggtgaac gagggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga 240
cgcttgctc gagcaggacc cagacagcaa agttgctgc gaaacatgca cc 292

<210> 2397
<211> 271
<212> nucleic acid
<213> Glycine max
<400> 2397

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ccccactaca accaccacac ctctcctacg ttacgctac cctttctgc tcttcttcta 120
cctttcaagt tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt 180
gaacgagggg caccctgaca agctctgcga ccaaactctc gatgctgtcc tcgacgcttg 240
cctcgagcag gaccagaca gcaaagttgc c 271

<210> 2398
<211> 287
<212> nucleic acid
<213> Glycine max
<400> 2398

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ttactgtctg ttcaagctac catctcgtc tcgctttctt agtgctcct tggccanaag 120
tnaaaatggc ccaagaaact tncctattca catctgaatc agtgaacgag gggcaccctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgtcttgga gcaggaccct 240
gacancaagg ttgcctgtga aactgcacca agaccaacat ggtgatg 287

<210> 2399
<211> 307
<212> nucleic acid
<213> Glycine max
<400> 2399

gcangcncgc gtncgtgagc tcggnttng gctcgnnggc caggcaagcn cactcancc 60
accacacctc tntcgttca cgtacccct ttctgctctt cttctacctt tcangtttta 120
anagtataaa gntggcagag acnttctnt ttncctcaga gtcggtgaac gagggacacc 180
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgttgcgctc gagnanggac 240
cnagacngca agttgcctgg gaaacatgca ccaggaccaa tttggtaatg gtctcgggaa 300
aatcgng 307

<210> 2400
<211> 291
<212> nucleic acid
<213> Glycine max
<400> 2400

ngtcgctgca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcaciaag 60
cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120
aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggnacc 180
ctganaagct cngngacnng nnntcngnng tngnncnng gngngctnga ggnggccct 240
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<210> 2401
<211> 304
<212> nucleic acid
<213> Glycine max
<400> 2401

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aaccaccaca cctctcctcg ttcacgtac cctttctgc tcttcttcta cctttcaagt 120

tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180
 caccctgaca agctctgcca cccaaatctc cgatgctgtc ctcgacgctt gncctcgagc 240
 aggaccaga cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcattgtct 300
 tcgg 304

<210> 2402
 <211> 302
 <212> nucleic acid
 <213> Glycine max
 <400> 2402

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 atgacccaag aatctttcct agtcacatct gaatcagcga acgagngcga cctgacaag 180
 ctccgtgacc agatctccga tgctgtgccc gatgcattct tggagcagga cccnacagca 240
 agtttgctg ttaaacctgt nccaagacca acatgggtgat ggtttgggaa anatcacaac 300
 cn 302

<210> 2403
 <211> 289
 <212> nucleic acid
 <213> Glycine max
 <400> 2403

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 tcctatttac ctcagagtcg gtgaacgn gn gacaccctga caagctctgc gaccaaactc 180
 ccgatgctgt ctcgacgct tgccctcgagc aggaccaga cagnaaagnt nccngcgaaa 240
 canntacca aaaccaacnt ngnnanngnc atnggagaaa ncacgacca 289

<210> 2404
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2404

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gagaagacga cagaaggggg cagcgcttga ttgaggccg ggcaagcccc actcaaccac 120
cacacctctc ctcggttcacg ctaccccttt ctgctcttct tctacctttc aagttttaaa 180
agtataaaga tggcagagac attcctatctt acctcagagt cgggtgaacga gggnnaccct 240
gacaagctct gcgaccaaatt ctccgatgct gtectcgacg cttgcctcga gcaggaccca 300
gacagcaaag ttgcct 316

<210> 2405
<211> 264
<212> nucleic acid
<213> Glycine max

<400> 2405
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tgaactaatc attaatctgc actcgctgtt tcagcttcat gcaccttct tttgcatcat 120
ttatatctct tgagaaatgg cacaagnac ctttctattc acatctgant ctgtaaacga 180
gggtcacccc gacatgctgt ncgaccagat ctctgatgca gtacttgatg cgtgccttga 240
acaggaccct gacagcaagg ttgn 264

<210> 2406
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 2406
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cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120
tataaagatg gcagagacat tcctatttac ctgagatcg gtgaacgagg gacacctga 180
caagctctgc gaccaaattc gccgatgctg tcctgcgacg cttgcctcga gcaggaccca 240
gacagcaaag ttgcctggcg aaacatgtac caaaaccaac ttggatcatg tnttcggaga 300
aatcacga 308

<210> 2407
<211> 331
<212> nucleic acid

<213> Glycine max

<400> 2407

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tcatatatct ctncctgtct tctcttcttt ctacctctca agtttttgaa gtataaagat 120
ggcagagaca ttcttattca cctcggagtc agtgaacgag ggacaccctg ataagctctg 180
cgaccaaate tcgatgtgtt cctcgacgct tgctctgaac aggaccagat cagccangtt 240
gcctgcgnaa acatgcacca agaccaattg gtcatggtct tcggagagat caccaccagg 300
gccangntga cnncaagatc gtgcgtnaca c 331

<210> 2408

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2408

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cncancgntc ctcggttnac gctaccctt tctnggtctt ctncctacctt tcangtttta 120
aaagtntaaa gatggnagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180
ctgagaagct ctgcgaccaa atctccgatg tgctctcgac gcttgccctg agcaggaccc 240
agacagcaaa gttgcctgcg aaacatgcac caaaaccaat tggnnaggtc ttcggagaat 300
caggacaagg ccaaggtgan t 321

<210> 2409

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 2409

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gntactgtct gttcaagcta ccctctctct ctctctttct tagtgccctc ttgccagaag 120
ttaaaatggc ccaagaaact ttcttattca catctgaatc agtgaacgag gggcaccctg 180
acaagctctg tgaccagatc tccgatgctg tgctcgatnc atgcttggag caaggccctg 240
ac 242

<400> 2410

Parameter	Unit	Value	Standard Error	95% CI	P-value
Intercept		1.00	0.00	1.00	0.00
Age	Year	0.02	0.01	-0.01, 0.05	0.15
Gender		0.05	0.02	-0.01, 0.11	0.08
Education	Year	0.01	0.00	0.00, 0.02	0.00
Income	Year	0.01	0.00	0.00, 0.02	0.00
Health		0.05	0.02	-0.01, 0.11	0.08
Marital Status		0.05	0.02	-0.01, 0.11	0.08
Occupation		0.05	0.02	-0.01, 0.11	0.08
Religion		0.05	0.02	-0.01, 0.11	0.08
Political Affiliation		0.05	0.02	-0.01, 0.11	0.08
Residence		0.05	0.02	-0.01, 0.11	0.08
Travel History		0.05	0.02	-0.01, 0.11	0.08
Exposure to COVID-19		0.05	0.02	-0.01, 0.11	0.08
Comorbidity		0.05	0.02	-0.01, 0.11	0.08
Healthcare Access		0.05	0.02	-0.01, 0.11	0.08
Healthcare Quality		0.05	0.02	-0.01, 0.11	0.08
Healthcare Cost		0.05	0.02	-0.01, 0.11	0.08
Healthcare Satisfaction		0.05	0.02	-0.01, 0.11	0.08
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00
Healthcare Quality (Ref)		1.00	0.00	1.00	0.00
Healthcare Cost (Ref)		1.00	0.00	1.00	0.00
Healthcare Satisfaction (Ref)		1.00	0.00	1.00	0.00
Healthcare Access (Ref)		1.00	0.00	1.00	0.00

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<212>	nucleic acid
<213>	Glycine max

<400> 2412

<210> 2413

<211> 121
 <212> nucleic acid
 <213> Glycine max

<400> 2413

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 cgagcctttg tctgtctttg ttgacaccta tggcaccggg aagatccatg ataaggagat 120
 t 121

<210> 2414
 <211> 273
 <212> nucleic acid
 <213> Glycine max

<400> 2414

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 gcttcttcag cttgagaaat ggcacaagaa acctttctat tcacactgaa tctgtaaacg 180
 agggtcaccc cgacaagctg tgcgaccaga tctctgatgc agtgcctgat gcgtagcctg 240
 aacaggaccc tgacagcaag gttgcctgtg aga 273

<210> 2415
 <211> 314
 <212> nucleic acid
 <213> Glycine max

<400> 2415

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 gagcgactga actaatcatt aatttgact cgctgtttca gttcatcac ccttgctttt 120
 gcatcattta tatctcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180
 taaacnaggg tnaccccgac angcnntnccg anccagatct ctgatgcagt acttgatgag 240
 tgccttgaac aggnccctga cngcaagggt cctggnnaga catgcaccag ggcnaacag 300
 ggtaagggnc ttgg 314

<210> 2416
 <211> 295
 <212> nucleic acid

2416
2417
2418
2419

<213> Glycine max

<400> 2416

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agtgaggcca ggcaagcccc actcaanac cacactttctc ctctttcacg ctaccccttt 120
actnctcttc ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt 180
tacctcagag tcggtgaacg agggacaccc tganaagctc tgcgaccana tctccgatgc 240
tgtcctcgac gttgcctcga gcaggaccca gacagcaaag ttgcctgcga aacat 295

<210> 2417

<211> 250

<212> nucleic acid

<213> Glycine max

<400> 2417

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ctctttctct acctttcatg ttttaaaagt ataaagatgg cagagacatt cctatttacc 120
tcagagtcgg tgaacgaggg acancctgac aagctctgcg accaaatctc cgatgctgtc 180
ctcgacgctt gcncgagcag gaccagaca gcaaagttgc ctgngaaca tgcnacaaaa 240
accaanttgg 250

<210> 2418

<211> 206

<212> nucleic acid

<213> Glycine max

<400> 2418

attaacttgg accttaagag ggggtggcat aggttctca agacagctgc ttanggacac 60
tttgaaggg atgatgcaga cttcacctgg gaagttgtga agccactcaa gtcagagaag 120
cctcaagctt aagagtgttg ttaagttaat cactcccttc agtggatgtc ttgctgggtg 180
tggatgaata atttgcgtgt ttcatg 206

<210> 2419

<211> 152

<212> nucleic acid

<213> Glycine max

<400> 2419

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ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 120

tcgatgcatg cttggannag gaccctgaca gt 152

<210> 2420

<211> 319

<212> nucleic acid

<213> Glycine max

<400> 2420

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gcagagncat tcctatttac ctcagantcn gtgaacgagg nacnccctgn cangetgctg 120

gtgaccaaat ctctgatgcn gtcctcgacg cttgcctcga acaggacnca nacancangg 180

ttgcctggng aancatgcac caaaaccaac ttgggtccatg gtcttcggag aaatcacgac 240

caaggccnat gttgactacg agaagatagt gcgtgncacc tgcagagcnt cggctttgtn 300

ctcaaacgat gtgggatgg 319

<210> 2421

<211> 262

<212> nucleic acid

<213> Glycine max

<400> 2421

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gctttcttag tgctccttg ccagaagtta aaatggcca agaaactttc ctattcacac 120

ctgaatcagt gnacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180

tcgatgttgc ttggagcagg accctgacag caaggttgcn ngnaannnct gcaccangac 240

aagnnnggtt ttgttgncag ac 262

<210> 2422

<211> 231

<212> nucleic acid

<213> Glycine max

<400> 2422

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gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120
agttaaaatg gcncaagaaa ctttctatt cacatctgaa tcagtgaacg aggggcaccc 180
tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcattgcttg a 231

<210> 2423
<211> 248
<212> nucleic acid
<213> Glycine max
<400> 2423

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tgaactaatc attaatattgc actcgctgtt tcagcttcat cacccttctt ttgcatcatt 120
tatatctctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180
ggtcaccccg acangctgtn cnaccagatc tctgatgcag tacttgatgc gtgccttgan 240
caggaccc 248

<210> 2424
<211> 322
<212> nucleic acid
<213> Glycine max
<400> 2424

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agcgggttac tgtctgttca agctaccatc tctctctctc ctttcttagt gcctccttgc 120
cagaagttaa aatggcccaa gaaactttcc nattcacatc tgaatcagtg aacgaggggc 180
accctgaaca agctctgtga ccagatnctc cgatggctgt gctcgatgnc atgcttggag 240
caggaccctg acagcaaggt tgctgtgaa acctgcacca ggaccaacat ggtgatgggt 300
ttcgagagat cacaaccaag gc 322

<210> 2425
<211> 317
<212> nucleic acid
<213> Glycine max
<400> 2425

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accaccacac ctctcctcgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180
accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240
accagaaca gcaaagttgc ctgcgaaact ggcaccaaca ccaattggtc atggtontcg 300
gagaaatcnc gaccagg 317

<210> 2426
<211> 287
<212> nucleic acid
<213> Glycine max

<400> 2426

angcacgcgt acgtaagctc ggaattcggc tcgagcggct gcgagaagac gacagaaggg 60
ggcagcgctt gatttgaggc caggcaagcc ccacttcaac caccacacct ctctcgttc 120
acgtacccc tttctgctct tcttctacct ttcaagtttt aaaagtataa agatggcaga 180
gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 240
aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagaca 287

<210> 2427
<211> 347
<212> nucleic acid
<213> Glycine max

<400> 2427

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gaagacgaca gaagggggca gcgcttgatt tgaggccagg caagccccac tcaaccacca 120
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 180
tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacacctga 240
caagctctgc gaccaaactt ccgatgctgt cctcgacgct tgccctcgagc aggaccaga 300
cagcaaattg cctgcgaaac atgcaacaaa aacaanttgt canggnc 347

<210> 2428
<211> 288
<212> nucleic acid

<213> Glycine max

<400> 2428

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tgtctgttca agctacncat ctctctctct cttncctagt gcctccttnc cagaagttan 120
natggcccaa gaaactttcc tattcacatc tgaatcagtg aancgagggg caccctgaca 180
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gangngacag 240
canggttgcc tgtgaaacct gcaccaagan caacatggtg atgntttt 288

<210> 2429

<211> 226

<212> nucleic acid

<213> Glycine max

<400> 2429

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cacgctaccc ctttctgctc ttntttctac ctttccaagt tttaaaagtn taaagatggc 120
agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca agctctgcga 180
ccaaatctcc gatgctgtcc tcgacgcttg cctcgagcag gaccca 226

<210> 2430

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2430

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gcagcgcttg atttgaggcc aggcaagccc cactcacacc accacacctc tcctcgttca 120
cgctacccct ttctgctctt cttctacctt tcaagtttta aaagtataaa gatggcagag 180
acattcctat ttacctcaga gtcggtgaac gagggacacc ctgacaagct ctgcgaccaa 240
atctccgatg ctgtcctcga cgcttgccctc gagcaggacc cagacag 287

<210> 2431

<211> 164

<212> nucleic acid

<213> Glycine max

<400> 2431

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 tcccagctc atgcccttga gccatgtcct tgccacgaag ctcggtgna agctcancga 120
 ggttcggaan aacggganat gcccttggnt gannccntnt ggca 164

<210> 2432

<211> 292

<212> nucleic acid

<213> Glycine max

<400> 2432

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 tgcaagctct gcttcacgag agtggtcttt ctctgtttca acaccttaat ttgcacacgc 120
 tgcttcttca gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180
 cgaggggtcac cccgacaagc nnncgaccag atngcnnang cagtgcgcga ngngngncnt 240
 nnacaggacc cngncagcaa ggcnngctgn nagacangca ncaagaccaa ca 292

<210> 2433

<211> 97

<212> nucleic acid

<213> Glycine max

<400> 2433

ccaagaccaa catggtcatg gtctctggag anctcacaac caaggccacc gtagactang 60
 agaagattgt ccgtgacaca tgccgcgaaa ntggata 97

<210> 2434

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 2434

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 accatctctc tctctctttc ntagtgtctc ttgccagaag ttaaaaatgg cccaagaaac 180
 tttcttattc acatctgaat cagtgaacga ggggcacctg acaagctctg tgaccagatc 240

tccgatgctg tgctcgatgc atgcttggag caggacctga cagcaagggt gctgtgaaac 300
 ctgcaccaan 310

<210> 2435
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2435

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 ctcgtncaag ctaccccttt ctgctcttct tctacctttc aagttttaaa agtataaaga 180
 tggcagagac attcctattt acctcagagt cggatgaacga gggacaccct gacaagctct 240
 ggcaccaaatt ctccgatgct gtctcncgac gcttgccctg ancaggaccc agacagcaaa 300
 gttgcc 306

<210> 2436
 <211> 278
 <212> nucleic acid
 <213> Glycine max
 <400> 2436

actnccgatg cagcgcgtacg taagctcgga attcggctcg agttgaggcc aggcaagccc 60
 cactcaacca ccacacctct cctcggtcac gctacccctt tctgctcttc ttctaccttt 120
 caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180
 agggacaccc tgacaagctc tgcgacaaa tctccgatgc tgcctcgcac gcttgccctg 240
 anggccaga cagcaagntt gcctgcgaaa catnccnc 278

<210> 2437
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 2437

gcacgcgtac gtaagctcgg aattcggctc gagatttctt cccatccttt cttccttcac 60
 cactttgaac ttgaacttag ttgggggact tggctgagtt agactgtnat gtttaaattg 120

tagtcatggt ggtgtttttg gctgtgaatt tgctcatatg tgctaattat gtgtttcttgt 180
 ttgatgttac tctacagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 240
 agtgaacgag gggcaccctg aacaagctct gtgaccagat ctccgatgct gtgctcgatg 300
 catgcttgga gcagg 315

<210> 2438
 <211> 121
 <212> nucleic acid
 <213> Glycine max
 <400> 2438

attaacttgg accttaagag ggggtggcat aggttcctca agacagctgc tnatggacac 60
 ttggaaggg atgatgcaga cttcacctgg gaagttgtga agccactcaa gtcagagang 120
 c 121

<210> 2439
 <211> 289
 <212> nucleic acid
 <213> Glycine max
 <400> 2439

gtcgcacgca cgcgtagta agctcggaat tcggctcgag caggcaagcc ccaactcaacc 60
 accacacctc tctcgttca cgctaccctt ttctgtcttt cttctacctt tcaagtttta 120
 aaagtataaa gatggcagag anattcctat ttacctcaga gtcggtgaac gagggacacc 180
 ctgacaagct ctgcgancca aatctccgat gctgtcctcg acgcttgctt cgagcaggnc 240
 ccagacagcc aaagttgcct gcgaaacang cagcnaaacc aacttggtc 289

<210> 2440
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2440

cntcangnac gcgtacgtaa gctcggaatt cgggctcgag nttgaggcca ggcaagcccc 60
 actcaaccac cacacctctc ctcgttcacg ctaccctttt ctgctcttct tctacctttc 120
 aagtttttaa agtataaaga tggcagagac attcctattt acctcagagt cgggtgaacga 180

gggacaccct gacaagctct ggggacaaa tctccgatgc tgcctcgcac gcttgccntn 240
gagcaggacc cnganagcaa antngcttgg gaaanttgcn caaaaacat ttgggtnnng 300
gtntgngnaa 310

<210> 2441
<211> 283
<212> nucleic acid
<213> Glycine max
<400> 2441

angcacngt acgttagctc ggaattcggc tcgagcttaa caacagcaca aagcgggtta 60
ctgtctgttc aagctaccat atctctctct ctnttcttag tgcctncctt gccanaagtt 120
aaaatggnc ntgaaacttt ccnattcaca tntnnatcag tgaacgaggg gcaccccgac 180
aagctctntn atagatcngg gtngncagtg ctagatgnat gnttggagca ngancctnan 240
agcnaggntn cctgtgaaac cnggcacna gaccaacatg gtn 283

<210> 2442
<211> 240
<212> nucleic acid
<213> Glycine max
<400> 2442

tgaggccagg caagcccccac tcaaccacca canatnanc ctcgttncac gctacccctt 60
ctcncctctt cttctacctt tcangtttta anagtataaa gatggcagag acattcctat 120
ttacctcaga gtcggtgaag agggacaccc tgacaagctc tgcgacaaa tctccgatgc 180
tgtcctcgcac gcttgctcgc agcaggaccc agacagcaaa gttgntggaa acatgcacca 240

<210> 2443
<211> 296
<212> nucleic acid
<213> Glycine max
<400> 2443

tcgatgcac gcgtacgtaa gctcggatt cggctcgcag gtttttgaag actgctgcct 60
atggacactt tggaagagaa gaccctgact tcacatggga agtgggtcaaa cccctcaagt 120
gggagaaggc ctaagtaant cattccactg ctctatgctg gaagtttttt gagcgttgcc 180

cttataatat gtctaataatc cataactttc cacgtctctt acnctgtgtg tttctctctt 240

cttctctcta ttttggttatt tgtatgttct tttgtaattt ttacgtgatc aactaa 296

<210> 2444

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2444

gtcgcangca cgcgtacgta agctcggaat tccgctcgag ngtggccata ggttctctca 60

gacagctgct tatggacact ttggaaggga tgaccctgac ttcacctggg aagttgtgaa 120

gccactcaag tctgagaagc ctcaagctta agattgttgt gaagttaatc actcccttca 180

atggatgtct tgctaggtgt ggatgaataa tttgcgtgtt ccatgactac tactacttca 240

ttcataggtc taatgtcatc tcatcaatac ttaaactggt ttttttt 287

<210> 2445

<211> 185

<212> nucleic acid

<213> Glycine max

<400> 2445

agcagactta acaacagcac aaagcgggtt actgtctgtt caagctacca nnnnnnnnnn 60

nnnnnnntag tgctctcttg ccagaagtta aaatggccca agaaactttc ctattcacat 120

ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180

togat 185

<210> 2446

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2446

ngtconatgca cgcgtacgta agctcggaat tccgctcgag gttnggttct gcacgctctg 60

cttcacagca gtgttctttc ttcgtttcaa caccttanat ttgcacacgc tgcttcttca 120

gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa cgaggggtcac 180

cccagacaagc tgtgcgacca gatctctgat gcagtgtctg atgcgtg 227

<210> 2447
 <211> 98
 <212> nucleic acid
 <213> Glycine max

<400> 2447

ccttttcagg gaaggaccct accaagggtg acagaagtgg tgcctatatt gtaaggcagg 60
 ctgcaaagag tgtcgtggca aatggccttg ctagaagg 98

<210> 2448
 <211> 304
 <212> nucleic acid
 <213> Glycine max

<400> 2448

gtcgngcac gcgtacgtna gctcggaatt nggctcgagc tcgagccgca acagcacaaa 60
 gggggttact gtctgttcaa gctaccatct ctctctcact ttcttngtgc ctcttgcca 120
 gangttanna tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180
 ncnnnacaag ctctgtgacc agatctccga tgctgtgcta cgatgcatgc ttggagcang 240
 naccctgaca gcnaagttgc ctgtgaaacc tggcaccaag ancaacatgg tgatggtttt 300
 cgga 304

<210> 2449
 <211> 266
 <212> nucleic acid
 <213> Glycine max

<400> 2449

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gtgatttga gtttgagcg 60
 actgaactaa tcnttanttt gactcgtg tttcagcttc atcacccttc ttttgcata 120
 tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180
 agggtcaccc cgacangctg tgcgaccaga tctctgatgc agtacttgnn gcgngcctna 240
 aaggncacca ncancaaggt cgcctg 266

<210> 2450
 <211> 159
 <212> nucleic acid

<213> Glycine max

<400> 2450

tccgaattcg gctcgagaac agcacaagc gggttactgt ctgttcaagc taccatctct 60
ctctctcttt cttagtgcct ccttgccaga agttaaaatg gccaagaaa ctttcctatt 120
cacatctgaa tcagtgaacg aggggcaccc tgacaagct 159

<210> 2451

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2451

gtcgcangen ngcgtnacgn nnagctcggn attcggtcg agccaagccc cactcaacca 60
ccanaccact ctctctgtct ttcttctacc tttccaagtt tntaaagtat taagntggca 120
gagacantcc nanntanctc agagncnng nangngggnc ancctgnan gcgctncgac 180
naatctnca tgtgtctctg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240
cgaaacatgc accaaganca attgggtcatg gtcttcggag agatcacca 289

<210> 2452

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2452

gtcgcattgca cgcgtacgtn agctcggaat tcgggtcgag ctcgagccgc tccacccaac 60
acgacgagac tgtcaccaat gangaaattg ntgntgacct tcaaanagca tgtgatcaan 120
cctgngatnc cngngaantn nctnatnagn agnncanttt ccnattngaa cnttaaggc 180
ggtttggaac tgggtggccn nnaggggcna ngctgggtct ccggggncga aaagancctt 240
atnccgatat ttagngngna nggggtgccc atgggggtgg tgnntcnccg ggan 294

<210> 2453

<211> 181

<212> nucleic acid

<213> Glycine max

<400> 2453

gnngetgcac gcgtacgtna gctcggaatt cggctcgagt aacaacagca caaagcgggt 60
 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180
 a 181

<210> 2454
 <211> 268
 <212> nucleic acid
 <213> Glycine max
 <400> 2454

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag tgggagttag gttctgncac 60
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120
 cttcagcttg aggaatggca caagaaacct ttctnttcac atctgantct gtanacgang 180
 gtcacccoga caagntgtgc gaaccagatn ctctgatggc agtgctcatg cgtgcgctga 240
 ncaggacct gacagcaagg ttgcctgn 268

<210> 2455
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2455

cntcgcangc acgcgtacgt gagctcgga ttcggctcga gcaggcaaan cccacncaac 60
 cgncanangg aaaaaacgaa ngcgcnaccc cattcngctc ntcttcnacc tnnrangann 120
 naagagnnta aagatggcag agacattcct atttacctca gagtacggtg aacgagggac 180
 accctgacaa gctctgcan ccaagtctcc gatgctgtcc tcgncgcttg cctcgagcag 240
 gaccagaca gcaaagttgc gcagcganac atgcancaag ncgnattggn ccatgggtt 298

<210> 2456
 <211> 154
 <212> nucleic acid
 <213> Glycine max
 <400> 2456

caccancagc aaacaannaa naagtcgcat gcacgcgtac gtaagctcgg aattcngctc 60

gaggggaaga cccaagtac cgttgagtat tacaatgaca atgggtgccag gggttcctatt 120
ccgtgtacac accgtgctaa tttccacaca acat 154

<210> 2457
<211> 284
<212> nucleic acid
<213> Glycine max

<400> 2457

cgtcgcangc acgcgtacgt aagctcgga ttcggctcga gcacaaagcg gggttactgtc 60
tggttcaagct accatctctc tctctctttc ttagtgccctc cttgccagaa gttaaaatgg 120
cccaagaaac tttctatttc acatctgaat cagtgaacga ggggcaccct gacaagctca 180
gagaccagaa nancgangcn gngcacgacg caagtccttg agcaggaccc agtacagcaa 240
ggnnnnncagt gnaaccngta ccaagaccaa caggtgatgg tcct 284

<210> 2458
<211> 213
<212> nucleic acid
<213> Glycine max

<400> 2458

gtcncangca cacgtacgta agctcggaat tcnnctcgag ncaagcccca ctcaaccacc 60
acacctctcc tegtacacgc tacccttttc tgctcttctt ctanctttca agtttttaaaa 120
gtataaagat ggcagagaca ttcctattta cctcagagtc ggtgaacgag ggacaccctg 180
acaagctctg cgaccaacan ctccgatgct gtc 213

<210> 2459
<211> 217
<212> nucleic acid
<213> Glycine max

<400> 2459

gtngcatgca cgcgtacgtn agctcggaat tcggctcgag caacagcaca aagcgggtta 60
ctgtctgttc aagctacctt ctntctctct ctttcttagt gcctccttgc cagaagttaa 120
aatggnccaa gaaacttttc tattcacatc tgnatcagtg ancgangggc accctgacaa 180
gctctgtgac cagatctccg atgctgtgct cgntgca 217

<210> 2460
 <211> 233
 <212> nucleic acid
 <213> Glycine max

 <400> 2460

 ctangcacgc gtacgtnagc tcggaattcg gctcgagcag caaaggagtg atttggagtt 60
 tggagcgact gaactaatca ttaatttgca cttcgctgtt tcagcttcat cacccttctt 120
 ttgcatcatt tataatctctt gagaaatggc acaagaaacc tttctattca catctgaatc 180
 tgtaaacgag ggtcaccccg acangctgtg cgaccagatc tctgangnag nat 233

<210> 2461
 <211> 202
 <212> nucleic acid
 <213> Glycine max

 <400> 2461

 gncgcangca cgcgtacgta agctcggaat tcggctcgag gggcaagccc cactcaacca 60
 ccacacctct cctcgttcac gctacccctt tctgctcttc ttctaccttt caagttttta 120
 aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180
 tgacaagctc tgcgaccaa tt 202

<210> 2462
 <211> 196
 <212> nucleic acid
 <213> Glycine max

 <400> 2462

 nccnnacgtc gcntgcacgc gtacgtaagc tcggaattcg gctcgaggca gacttaacaa 60
 cagcacaaaag cgggttactg tctgttcaag ctaccatnct ctctctctct ttacttagtg 120
 cctccttgcc agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga 180
 acgaggggca ccctga 196

<210> 2463
 <211> 323
 <212> nucleic acid
 <213> Glycine max

<400> 2463

gtcgcangca cgcgtacgta agctcgggaat tcggctcgag ttaacaacag cacaaagcgg 60

gttactgtct gttcaagcta ccattctctct ctctctttct tagtgcctcc ttgccagggtg 120

ctgccactct ctctttctct ctcttcatcc ttctgttggg ttggttgtgg agtgttgttt 180

tctgttgtgc acgtgttgtc attttttacc ctgccacag atctgaagtg ttcaagtttg 240

gattttgtgc ttctggaagt taaaatggcc caagaaactt tcctattcac atctgaatca 300

gtgaacgagg ggcaccctga caa 323

<210> 2464

<211> 132

<212> nucleic acid

<213> Glycine max

<400> 2464

gcatgcacgc gtacgtaagc tcggaattcg gctcgagcag atggcaagac acaagtaact 60

gttgagtaac tacaatgaca atggtgccat ggttccagtt cgtgtccaca ctgtcctaata 120

ttccacacac ac 132

<210> 2465

<211> 189

<212> nucleic acid

<213> Glycine max

<400> 2465

gggagttagg ttctgcacgc nctgnngcca gcgagtgtgc tntcttcgtg tcaacacctg 60

aatttgcann acgctgcgnc tgcagcttga gaaatggcac aagaaaccnn gctatncana 120

tctgaatcgg taaacgaggg tcacnncgac aagctgtggg accagatctc tgatgcagtg 180

ctcgatgcg 189

<210> 2466

<211> 138

<212> nucleic acid

<213> Glycine max

<400> 2466

tngcacgcgt acgtaagctc ggnattcggc tcgagggagn ttggtgntgg tgaccaaggt 60

catatgttcg gctatgccna ctgacgagnn ctcccagagct catgcccttg agccatgtnc 120
 ottgccacga agcttcgg 138

<210> 2467
 <211> 341
 <212> nucleic acid
 <213> Glycine max
 <400> 2467

ncctctnanc ntttacatcg anacnacgta cgtnacgtcg gnattcggct cgagcagact 60
 taacaacagc acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt 120
 agtgccctcc ttgccaggtg ctgccactct ctctttctct ctcttcatcc ttctgttggg 180
 ttggttgttg agtggtgttt tctgttgtgc acgtgttgtc attttttacc ctgccacag 240
 atctgaagtg ttcaagtttg gattttgtgc ttctggaagt taaaatggcc caagaaactt 300
 tcctattcac atctgaatca gtgaacgagg ggcacctga c 341

<210> 2468
 <211> 273
 <212> nucleic acid
 <213> Glycine max
 <400> 2468

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag aatgacacca cctacccct 60
 tctccctata aatggcaact caatgcccc cttagaactc gcagcgcttg atttgaggcc 120
 aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt tctgctcttc 180
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 240
 tcggtgaacg agggacaccc tgacaagctc tgc 273

<210> 2469
 <211> 181
 <212> nucleic acid
 <213> Glycine max
 <400> 2469

ggcgcacnca gnnttcggct cgagggccag gcaagcccca ctcaaccacc acacctctcc 60
 tcgttcacgc tacccttttc tgctcttctt ctacctttca agttttaaaa gtataaagat 120

ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg acaagctctg 180

c 181

<210> 2470

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 2470

gtgcgatgca cgcgtacgta agctcggaat tcggctcgan gnagacttaa cancagnnca 60

aagcgggtta ctgtctgntc angctaccat ctctcnctct nttcttagt gcctccntgc 120

cagnagttnn aatggcccaa gnnacttten tantcacatc tgantcnntg aacgaggggc 180

accngataa gctctgtgan cagatctccg atgctgtgct ccgatgnatg cttggagcng 240

gnnnctgnca gcnaggntgn ctgtgnaach tgcacnangn ncancatggt gatggntttc 300

ggnga 305

<210> 2471

<211> 199

<212> nucleic acid

<213> Glycine max

<400> 2471

agtcgangca ncgtacgtaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60

agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120

agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180

ccctgacaag ctctgtgat 199

<210> 2472

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2472

cgcatgcacg cgtacgtaag ctcggaattc ggctcgagcc ttgatctcaa gaggggtgga 60

aatggcaggt tcttgaagac tgctgcatat ggacactttg gcagagatga ccctgacttc 120

acatgggaag tggatgaagcc actcaagggg gagaaggtag ctgcttaact aaaaggggtt 180

ccaacactct tggcaaggga cttttgcact actactggct tcttattatc tgattgctaa 240
aattttctct atgtttcctt cctctact caattctgtt ttttttttnc ngatattttt 300
tatgaatttc cccctttttt ttgtgta 327

<210> 2473
<211> 256
<212> nucleic acid
<213> Glycine max
<400> 2473

ggggtggaaa tggcagggtc ttgaagactg cngcatatgg acactttggc agagatgacc 60
ctgacttcac atgggaagtg gtgaagccac tcaaggggga gaaggtagct gcttaactaa 120
aaggggttcc aacactcttg gcaagggact tttgcactac tactggcttc ttattatctg 180
attgcnaaaa tttnctctat gnntccttcc ctcttactca attctgtttt tntttttctg 240
tatttttnat gaattc 256

<210> 2474
<211> 214
<212> nucleic acid
<213> Glycine max
<400> 2474

atggcagggtt cttgaagact gctgcatatg gacactttgg cagagatgac cctgacttca 60
catgggaagt ggtgaagcca ctcaaggggg agaaggtacc tgcttaacta aaaggggttc 120
caacactctt ggcaagggaac ttttgcacta ctactggctt cttattatct gattgctaaa 180
attttctcta tgtttccttc cctcttactc aatc 214

<210> 2475
<211> 206
<212> nucleic acid
<213> Glycine max
<400> 2475

atggcagggtt cttgaagact gctgcatatg gacactttgg cagagatgac cctgacttca 60
catgggaagt ggtgaagcca ctcaaggggg agaaggtacc tgcttaacta aaaggggttc 120
caacactctt ggcaagggaac ttttgcacta ctactggctt ctattatctg attgctaaaa 180

tttttctctat gtttccttcc ctctta

206

<210> 2476

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2476

cctccncaac agtcgcatgc acgcgtacgt aagtcggaat tcnctcgag ctgctgcata 60

tggacacttt ggcagagatg accctgactt cacatgggaa gtggtgaagc cacttcaagg 120

gggagaaggt acctgcttaa ctaaaagggg ttccaacact cttggcaagg gacttttgca 180

ctactactgg cttcttatta tctgattgct aaaattttct ctatgtttcc tccctcttac 240

tcaattctgt ttttttttnt ctgtnttttc tnatgaattt cccctttttt tttgggnact 300

ngnatgtgtt c 311

<210> 2477

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2477

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natgacctg acttcacatg ggaagtgggtg aagccacttc aagggggaga aggtacctgc 120

ttaactaaaa ggggttccaa cactcttggc aagggaactt tgactacta ctggcttctt 180

attatctgat tgctanaatt ttctctatgt ttctttccct ctactcaat tctntttttc 240

nttttctgta tttntttatg aatttcccc tttntntgn gnacttgtn gngtnctnnc 300

<210> 2478

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2478

acgtcgcatg cacgcgtacg taagctcggn attcggtcg agatttcgca tctnctcctt 60

ctcattccaa ctccaaaata cacacacacg atggaaacct tcctcttcac ctcagaatct 120

ncaaacgagg gccaccccgga caagatctgt gaccaggttt ctgatgccat cctcgatgct 180

tgcttgagc aagaccaga gagcaaggtt gcctgcgaga cctgtacaaa aaccaacatg 240
gtcatggtct ttggggagat cacaaccaag gccaaagtga actacgagaa t 291

<210> 2479
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 2479

agtcgatgc acgcgtacgt aagctcggaa ttcggtcga gcggctgcga gaagacgaca 60
gaaggggttc ctcttcaatt tcgcatcttc tcttctcat tccaacttcc aaaatacaca 120
cacacgatgg aaaccttctt cttcacctca gaatctgtaa acgagggcca ccccgacaag 180
atctgtgacc aggtttctga tgccatcttc gatgcttgc tggagcaaga cccagagagc 240
aaggttgctt gcgagacctg tacaaaaacc aacatggtca tggctcttgg ggagatcaca 300
accaaggc 308

<210> 2480
<211> 262
<212> nucleic acid
<213> Glycine max

<400> 2480

cttctgtttc tggnttcatt ngggtgtttg gacacaattg ctnaattata ctttctgttg 60
attgtgttga cgcnggactg aatgaactaa tggagtctan aggtgggaaa nagaagtcnn 120
nnnnnnnnnn nnnnnaatca ttgttctacg aagctccctt cggatacagc atnngaagac 180
gttagacca aaggtggaat caagaaattc agatctgctg cttactccaa cgtatatntt 240
cttctgatgc agtgattctg ta 262

<210> 2481
<211> 420
<212> nucleic acid
<213> Glycine max

<400> 2481

ggnggttnng aatgatttag gtgnoctagc caagacanat gacatcgctg ncacgcntac 60
gtaagntcgg aattcggctc gagctcgagc cgcaaaaatt cgggaagact agtgactgtg 120

gttcacctag atcaactcta aagtgtctga atgaggaaga tgaggaagag tagtttcctt 180
aagtgtcttt attattctgt ccttgtgaaa ataagtctgg ttttccagat acgttattgt 240
ttttctttgt tgtctttttt agcttctgtt agagaccatt tgggcattta gacctttatt 300
gtttctatta ccatttgaac atcgaatgga ttaataaatc actttgtttg cgtgcaaaaa 360
aaannacana tctttcnana aanaaaaaaa annaaaaana acanaaaaaan aaaaaaaaaan 420

<210> 2482
<211> 287
<212> nucleic acid
<213> Glycine max
<400> 2482

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc nggaggatta ggccctccgag 60
cattgtccaa agaccaattg gacgagattc ttaaaccagc agagtgcact attgtngcat 120
cactttcaaa tgattatgtt gactcttatg ttctgtcaga gtcaagcctg ttctgtctatc 180
cttataaaat tatcatcaaa acttgtggga ctaccaaatt gcttctgtcc atccctgtcc 240
attctcaagt tgggctgatg ctcttgacat agctgtgaaa tctgtga 287

<210> 2483
<211> 288
<212> nucleic acid
<213> Glycine max
<400> 2483

acgtcgcang cacgcgtacg taagctcgga attcggctcg agcgaaaaaa gacttgagat 60
atcatttttt gaaaatgggtg tgtttgctga ccccgaggga ttaggcctcc gagcattgtn 120
ccaaagacca attggacgag attcttaaac cagcagagtg cactattgtt gcatcacttt 180
caaagatta tgttgactct tatgttctgt cagagtcaag cctgttcgtc taccctataa 240
aattatcatc aaaacttgtg ggactaccaa attgcttctg tccatccc 288

<210> 2484
<211> 306
<212> nucleic acid
<213> Glycine max
<400> 2484

cgctcangca cgcgtagcta agctcggaat tcggctcgag ntctgatata tctgantttg 60
 antttgaccc ntgcggatat tcaatgaatg gnntagaagg gagtgctata tccaccatcc 120
 atgtcactcc tgaagatggg ttcagttacg caagttttga agctgttggt tatgacttta 180
 atgacatggc tctaggtgaa nttgtggaaa gganttttagc ttgcttttgt ccagcagagt 240
 tttctgttgc tttgcacatt gacatgcatg gtgagaaaact aanaaatttc ccttaganat 300
 caaagg 306

<210> 2485
 <211> 314
 <212> nucleic acid
 <213> Glycine max

<400> 2485

gtctatgcac gcgtacgtaa gcnncggaat ncggctcgag gtctgatata tctgattttg 60
 aatttgaccc ttgcggatat tcaatgaatg gaatagaagg gagtgctata tccaccatcc 120
 atgtcactcc tgaagatggg ttcagttacg caagttttga agctgttggt tatgacttta 180
 atgacatggc tctaggtgaa cttgtggaaa ggatttttagc tgcttttgtc cagcagagtt 240
 ttctgttgc tgcacatgac atgcagnctg agaaantaat aaattcccta gacatcaaag 300
 gatactactg tggg 314

<210> 2486
 <211> 476
 <212> nucleic acid
 <213> Glycine max

<400> 2486

aactccnctc cgntcgnncg tgnnntngta cagaagtccc ggctcgaccc acacgtcagc 60
 tccttcaaac tccatctttc caaatcctct ctttgcgatt gtgttttgat ctgcttccta 120
 ctgcgatagt ttctctactg ttacatggcc atggcggttt ccgcaattgg ttttgaaggt 180
 ttcgagaaaa ggttggaaat atcctttttc cagccgggac tttttgctga ccttgaagga 240
 aggggtctaa gggctcttac aaaatcccag ttgggtgaga ttctaacc accagctgttc 300
 accattgttt cttegtctaa aaacgataat gtcgactcct atgttctatc tgagtccagc 360
 ctctttgttt atgcctacaa gatcatcatc aaaacctgtg gtactactaa gctattgctt 420

gcaatccac ccatattgaa gttcgtgaa atgctttccc ntaatgttaa gtongt 476

<210> 2487
 <211> 510
 <212> nucleic acid
 <213> Glycine max
 <400> 2487

gnngnaaggt tgntnagnaa cgggggtana tngaannnnn nnnnnnnnnn nnnnnnnnnn 60
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnngt gcatgactgg 120
 cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag ctgccatgat 180
 gactgttaat tccggcatta gaaaaattct tccagattcc gagatttggtg actttgactt 240
 tgaaccatgt ggttattcaa tgaactctgt tgaaggcgct gctgtttcta cgattcatgt 300
 taccocagaa gatgggttca gttatgcaag cttcgaaact gttggttatg acttcaaagc 360
 ggtgaatctg aacgaaatgg gttcaagang gtattggcat gtttcctncc aactgagttt 420
 ctgttgcatg caatgtggat ggtgcaagca agtcgtttga ccanacctgc ttctggatgt 480
 taagggatac tgnctgaaaa gaggaacccc 510

<210> 2488
 <211> 560
 <212> nucleic acid
 <213> Glycine max
 <400> 2488

gnnnngnnnn nnntnnnaag taagtnaaan ctctntggaa agcncctacc ggtncggaat 60
 tcccgccccg acccagcgt cgggttcag ctgccattat gactgttaat tctggcatta 120
 gaaaaattct tccagantct gagatttggtg actttgactt tgagccatgt ggttattcaa 180
 tgaactctgt tgaagggtgt gctgtttcta ccattcatgt taccocagaa gatggtttca 240
 gttatgcaag ctttgagact gttgggnatg acttcaaagt ggtgaatctg aacgaaatgg 300
 ttaagagggg attggcatgt tttctcccaa ctgagttctc tgttgcatgt catgtggatg 360
 gtgcaagcaa gttgtttgat cagacgtgtn ttctggatgt taagggatac tgtcgcaag 420
 agaggagccc acgaaagggg ttgggaatgg gtggnnntct tggctaccaa aaaaantgcc 480
 aaagacttgg gaactggggg tcaactagan ccaactctga aangntggaa aagaaggaag 540

atgaagaaag agtagttttt

560

<210> 2489
<211> 485
<212> nucleic acid
<213> Glycine max
<400> 2489

gngnnnnngn nnnnnnnnnn tnnnttttng tggcntanag cccnnnataa cancatcacg 60
tcgcangcat gannncgnaa ntcggctccg anggttattc aatgaactct gttgaaggtg 120
ctgctgtttc taccattcat gttaccccag aagatgggtt cagttatgca agctttgaga 180
ctgttgggta tgacttcaaa gtggtgaatc tgaacgaaat ggtaagagg gtattggcat 240
gttttctccc aactgagttc tctgttgagc ttcattgtga tgggtgcaagc aagttgtttg 300
atcagacgtg ttttctggga tgttaaggga tactgtcgcg aagagaggag ccacgaaggg 360
cttggaatgg gtggttctct tgtctaccaa aaatttgcca agacttgtga ctgtggttca 420
nctanattaa ctctgaaat gctgggaaag aaggaanatt aaanaanaat aattttcctt 480
aagtg 485

<210> 2490
<211> 339
<212> nucleic acid
<213> Glycine max
<400> 2490

gcatgactgg cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag 60
ctgccatgat gactgttaat tccggcatta gaaaaattct tccagattcc gagatttgtg 120
actttgactt tgaaccatgt gggtattcaa tgaactctgt tgaaggcgct gctgtttcta 180
cgattcatgt taccocagaa gatggtttca gttatgcaag cttcgaaact gttggttatg 240
acttcaaagc ggtgaatctg aacgaaatgg ttcagagggt attggcatgt ttcctcccaa 300
ctgagttttc tgttgacgtt catgtggatg gtgcaagca 339

<210> 2491
<211> 412
<212> nucleic acid
<213> Glycine max

<400> 2491

gtttcagtta tgcaagcttt gagactgttg ggtatgactt caaagtgggtg aatctgaacg 60
aatgggttaa gaggggtattg gcatgttttc tcccaactga gttctctgtt gcagttcatg 120
tggatgggtgc aagcaagttg tttgatcaga cgtgttttct ggatgttaag ggatactgtc 180
gcgaagagag gagccacgaa gggcttgga tgggtgggtc tcttgtctac caaaaatttg 240
ccaagaacttg tgactgtggt tcacctagat caactctgaa gtgctggaaa gaggaagatg 300
aagaagagta gttttcttaa gtgtctttat tatgtccttg cgaaaataag tccggttttc 360
cagacagtga ttgtttntct ttggtgnttt ttnccttnta tgtagacca tg 412

<210> 2492

<211> 504

<212> nucleic acid

<213> Glycine max

<400> 2492

agtgtntntg nnagggggga nnatggtaac actctcgaag actatgacgt cgcattgcacg 60
cgtacgtaag ctcggaattc ggctcgagt catgcaacca tagttttatt aggatttttt 120
cttctttgtt ttcaattagg tttttttgtt gctctccttc aaactccatc tttccaaatc 180
ctctctttgc gattgtgttt tgatctgctt cctactgcga tagtttctct actgttacat 240
ggccatggcg gtttccgcaa ttgggttttga aggtttcgag aaaaggttgg aaatatcctt 300
tttccagccg ggactttttg ctgaccctga aggaaggggt ctaagggctc ttacaaaatc 360
ccagttgggt gagattctaa caccagctgc ttgcaccatt gtttcttcgc tcaaaaacga 420
taatgtcgac tctatgttc tatctgagtc cagcctcttt ggttatgcct acaagatcat 480
catcaaaacc tgnnggacta ctaa 504

<210> 2493

<211> 347

<212> nucleic acid

<213> Glycine max

<400> 2493

tgcaattggt tttgaagggt tcgagaagan gctggaaata tcctttttcc agcngggact 60
ttttgctgac cctgagggaa tgggtttaag agctcttgca nagtcccagt tggatgagat 120

acttacaccg gctgcttgca ccattgtttc atctctcaga aatgatcatg tcgactccta 180
 tgttctgtct gagtccagtc tctttgttta tgcctacaag atcatcatca aaacctgtgg 240
 tactacaaag ctactgcttg caatcccacc catattgaaa tttgctgaaa tgctttcctc 300
 aatgtagatc tgtgnaatac accaggggaag ttcanctttt ccggtgt 347

<210> 2494
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 2494

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gccatggcgg tttctgcaat 60
 tggttttgaa ggtttcgaga agangctgga aatatccttt ttccagccgg gactttttgc 120
 tgacctgag ggaatgggtt taagagctct tgcaaagtcc cagttggatg agatacttac 180
 accggctgct tgcaccattg tttcatctcn cagaaatgat catgtcgact cctatgttct 240
 gtctgagtc agtctctttg tttatgcta caagatcatc atcaaaacct gtggtactac 300
 aaagctactg cttg 314

<210> 2495
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 2495

aangcacgcg tacgtaagct cggaattcgg ctcgagaaac aatctgcttc agctgccatt 60
 atgactgtta attctggcat tagaaaaatt cttccagatt ctgagatttg tgactttgac 120
 tttgagccat gtggttattc aatgaactct gttgaagggtg ctgctgtttc taccattcat 180
 gttaccccag aagatgggtt cagttatgca agctttgaga ctgttgggta tgacttcaaa 240
 gtggtgaatc tgaacgaaat ggtaagagg gtattggcat gttttctccc aactgagttc 300
 tctgttgcaag ttca 314

<210> 2496
 <211> 320
 <212> nucleic acid
 <213> Glycine max

<400> 2496

gtacgtanag ctcggaattc ggctcgagtg gcattagaaa aattcttcca gattctgaga 60
tttgtgactt tgactttgag ccatgtgggtt attccaatga actctgttga aggtgctgct 120
gtttctacca ttcatgttac ccagagaagat ggtttcagtt atgcaagctt tgagactgtt 180
gggtatgact caaagtgggtg aatctgaacg aaatgggttaa gaggggtattg gcatgttttc 240
tcccaactga gttctctgtt gcagttcatg tggatgggtgc aagcaagttg tttgatcaga 300
cgtgttttct ggatgttaag 320

<210> 2497

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2497

gtngcangca cgcgtacgta agctcggaat tcggctcgag gccatggcgg tttccgcaat 60
tggttttgaa ggtttcgaga aaaggttgga aatatccttt ttccagccgg gactttttgc 120
tgacctgaa ggaaggggtc taagggctct taaaaatcc cagttgggtg agattctaac 180
accagctgct tgcaccattg tttcttcgct caaaaacgat aatgtogact cctatgttct 240
atctgagtc agcctctttg tttatgccta caagatcatc atcaaaacct gtg 293

<210> 2498

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2498

nnntaanang tcgnangcac gcgtacgtaa gctcngaatt cggctcgagc ttcaaagtgg 60
tgaatctgaa cgaaatgggtt aagaggggtat tnnctgttt tctcccaact gagttctctg 120
ttgccgttca tgtgnatggg gcaagcaagt tgtttgatca gacgtgtttt ctggatgtta 180
agggatactg tcgcgaagag aggagccacg aagggttgg aatgggtggg tctcttgtct 240
accaaaaatt tgccaagact tgtgactgtg gttcacctag atcaactctg aagtgtctgga 300
aagaggaaga tgaagaagag tagtttt 327

<210> 2499

[illegible]

<210> 2502

<211> 315
 <212> nucleic acid
 <213> Glycine max

 <400> 2502

 ntgcgatgca cgcntacgta agctcggaat tcggctcgag tngcggtttc cgcaattggg 60
 tttgaagggtt tcgagaaaag gttggaaata tcctttttcc agccgggact ttttgctgac 120
 cctgaaggaa ggggtctaag ggctcttaca aaatcccagt tgggtgagat tctaacacca 180
 gotgottgca ccattgtttc ttcgctcaaa aacgataatg tcgactccta tgttctatct 240
 gagtccagcc tctttgttta tgcttacaag atcatcatca aaacctgtgg taactactaa 300
 gotattgctt gcaat 315

<210> 2503
 <211> 312
 <212> nucleic acid
 <213> Glycine max

 <400> 2503

 cangcacggg tacgtaagct cggaattcgg ctcgaggta attctggcat tagaaaaatt 60
 cttccagatt ctgagatttg tgactttgac tttgagccat gtggttattc aatgaactct 120
 gttgaagggtg ctgctgtttc taccattcat gttacccag aagatggttt cagttatgaa 180
 gctttgagac tgttgggtat gacttcaaag tgggtgaatct gaacgaaatg gttaagaggg 240
 tattggcatg ttttctccca actgagttct ctgttgacgt tcatgtggat ggtgcaagca 300
 agtngtttga tc 312

<210> 2504
 <211> 440
 <212> nucleic acid
 <213> Glycine max

 <400> 2504

 ccacgcgtcc gccagattga tagttcattg catgcaacca tagttttatt aggttttttc 60
 ttctttgnnt tcaattaggt ttttgctgct ctcttcaaa ctccgtcttt cogaatctc 120
 tctttgtgat tgtgttctgt tctgcttct accgcgatag tttctcttct gaagcatggc 180
 catggcgggtt tctgcaattg gttttgaagg tttcgagaag aggctggaaa tatccttttt 240

ccagccggga ctttttgctg accctgaggg aatgggttta agagctcttg caaagtccca 300
 gttggatgag atacttacac cggctgcttg caccattggt tcatctctca gaaatgatca 360
 tgtcgactcc tatggctctg ctgaagtcca gtctctttgg ttatgcctac aagatcatta 420
 tcaaaacctg ggttactaca 440

<210> 2505
 <211> 287
 <212> nucleic acid
 <213> Glycine max
 <400> 2505

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag ggcggtttcc gcaattgggt 60
 ttgaagggtt cgagaaaagg ttggaaatat cttttttcca gccgggactt tttgctgacc 120
 ctgaaggaag gggctctaagg gctcttaca aatcccagtt ggtgagatt ctaacaccag 180
 ctgcttgca cattgtttct tcgctcaaaa acgataatgt cgactcctat gttctatctg 240
 agtcacgct ctttgtttat gctacaaga tcatcatcaa aacctgt 287

<210> 2506
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2506

nngcatgcac ncgtacgtaa gctcggaatt cggctcgagg ctgtttctac gnttcatggt 60
 accccagaag atggtttcag ttatgcaagc ttcgaaactg ttggttatga cttcaaagcg 120
 gtgaatctga acgaaatngt tcagagggta ttggcatggt tcctcccaac tgagttttct 180
 gttgcagttc atgtggatgg tgcaagcaag tcgtttgagc agacctgctt tctggatggt 240
 aagggatact gtcgtgaaga gaggagccac gaagggcttg gaatgggtgg ttctgttg 298

<210> 2507
 <211> 505
 <212> nucleic acid
 <213> Glycine max
 <400> 2507

ccacgcgtcc gcttgatggc tactttggca aacttggtgc aggaagcaat gcttatattt 60

tgggtggcca anacnaagca cagaactggc atgtctactc tgcttctgca gattctgtaa 120
 ctcaatgtga caatgtttac actcttgaga tgtgcatgac tggcctggat agagagaaaag 180
 cacaggtttt ctacaaagaa caatctgctt cagctgccat gatgactgnt aattccggca 240
 ttagaaaaat tcttccaaat tcccagaatt gngactttgn ntttgaacca tgtggntatt 300
 caatgaactn tgnttgaaag gcncttggtg ttctacgatt catgggtancc ccagaagatg 360
 ggtcanntat tgcaagcttt gnaaactntt gggatgact ttaaagccgg ngaatntgaa 420
 cccaaaaggn ttaaanggat ttggcatggt tctccaact taantttctg tncaantcat 480
 tggggaangt gcaagcaagn ntttt 505

<210> 2508
 <211> 294
 <212> nucleic acid
 <213> Glycine max

<400> 2508
 gtgcgatgca cgcgtacgta agctcggaa tccgctcgag ctttgaacca tgtggttatt 60
 caatgaactc tgttgaaggc gctgntgttt ctacgattca tgttacccca gaagatgggt 120
 tcagttatgc aagcttcgaa actgttggtt atgacttcaa agcggatgaat ctgaacgaaa 180
 tggttcagag ggtattggca tgtttcctcc caactgagtt ttctgttgca gttcatgtgg 240
 atggtgcaag caagtcgttt gagcagacct gctttctgga tgtaaggga tact 294

<210> 2509
 <211> 296
 <212> nucleic acid
 <213> Glycine max

<400> 2509
 agtcgcnngc acgcgtacgt aagctcggaa ttcggctcga ggttttctac aaagaacaat 60
 ctgcttcagc tgccattatg actgttaatt ctggcattag aaaaattctt ccagattctg 120
 agatttgtga ctttgacttt gagccatgtg gttattcaat gaactctgtt gaagggtgctg 180
 ctgtttctac cattcatgtt accccagaag atggtttcag ttatgcaagc tttgagactg 240
 ttgggtatga cttcaaagtg gtgaatctga acgaaatggt taagagggtg ttggca 296

<210> 2510

<211> 254
 <212> nucleic acid
 <213> Glycine max

 <400> 2510

 ggactttttg ctgaccctga aggaaggggt ctaagggtc ttacaaaatc ccagttgggt 60
 gagattctaa caccagctgc ttgcaccatt gtttcttcgc tcaaaaacga taatgtcgac 120
 tectatgttc tatctgagtc cagcctcttt gtttatgcct acaagatcat catcaaaacc 180
 tgtgggtacta ctaagctatt gcttgcaatc ccacccatat tgaagttcgc tgaaatgctt 240
 tcccttaatg ttaa 254

<210> 2511
 <211> 299
 <212> nucleic acid
 <213> Glycine max

 <400> 2511

 tcgcatgcac gcgtacgtaa gctcggaatt cggtcgcaga ggcgctgctg tttctacgat 60
 tcatgttacc ccagaagatg gtttcagtta tgcaagcttc gaaactgttg gttatgactt 120
 caaagcgggtg aatctgaacg aaatggttca gaggggtattg gcatgtttcc tcccaactga 180
 gttttctgtt gcagttcatg tggatgggtgc aagcaagtcg tttgagcaga cctgctttct 240
 ggatgttaag ggatactgtc gtgaagagag gagccacgaa gggcttgga tgggtgggt 299

<210> 2512
 <211> 257
 <212> nucleic acid
 <213> Glycine max

 <400> 2512

 gtcgactcct atgttctatc tgagtccagc ctctttgttt atgcctacaa gatcatcatc 60
 aaaacctgtg gtactactaa gctattgctt gcaatccacc catattgaag ttcgctgaaa 120
 tgctttccct taatgttaag tctgtgaatt acaccagggg aagtttcatt ttccccagt 180
 ctcagccata tccccatgc aacttttctg aggaagttgc tattcttgat ggctactttg 240
 gcaaacttgg tgcagga 257

<210> 2513

<211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2513

 gtcgengcac gcgtacgtaa gctcggaatt cggctcgagc tcgagccgct cctatgttct 60
 atctgagtc agcctctttg tttatncta caagatcatc anccaaaacc tgtggtacta 120
 ctaagctatt gcttgcaatc ccaccatat tgaagtgcg tgaaatgctt tcccttaatg 180
 ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgcgcag ccatatcccc 240
 atcgcaactt ttctgaggaa gttgctattc ttgatggcta ctttgcaaa cttggtgcag 300
 gaagcaatgc 310

<210> 2514
 <211> 322
 <212> nucleic acid
 <213> Glycine max

 <400> 2514

 tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc agagggtatt ggcattgttc 60
 ctcccaactg agttttctgt tgcagttcat gtggatgggt caagcaagtc gtttgagcag 120
 acctgctttc tggatgttaa gggatactgt cgtgaagaga ggagccacga agggcttgga 180
 atgggtggtt ctgttgtcta ccaaaaattc gggaagacta gtgactgtgg ttcacctaga 240
 tcaactctaa agtgctggaa tgaggaagat gaggaagagt agtttcctta agtgtcttta 300
 ttattctgtc cttgtgaaaa ta 322

<210> 2515
 <211> 314
 <212> nucleic acid
 <213> Glycine max

 <400> 2515

 aannnngaen cgcangcncg cgnacgtnan ctcggaattn ngctcnaggt ctaaggntc 60
 ttacaaaatc ccagttgggt gagattctna caccagctgc ttgcaccatt gtttcttcgc 120
 tcaaaaacga taatgtcgac tcctntgttc tatctganc cagcctcttt gtntatgcct 180
 acnagatcat catcaaaacc tgtggtacta ctaagctatt gcttgcnato ccaccatat 240

tgaagttngc tganatgctt tcccttaatg ttaagtctgt gaattacacc aggggaagtt 300
tcattttccc cagt 314

<210> 2516
<211> 283
<212> nucleic acid
<213> Glycine max
<400> 2516

gatgcacgcg tacgtaagct cggaattcgg ctcgagggtga ctttgacttt gagccatgtg 60
gttattcaat gaactctgtt gaagggtgctg ctgtttctac cattcatgtt accccagaag 120
atggtttcag ttatgcaagc ttgagactg ttgggtatga cttcaaagtg gtgaatctga 180
acgaaatggg taagagggga ttggcatgtt ttctcccaac tgagttctct gttgcagttc 240
atgtggatgg tgcaagcaag ttgtttgatc agacgtgttt tct 283

<210> 2517
<211> 247
<212> nucleic acid
<213> Glycine max
<400> 2517

gtccagcctc tttgtttatg cctacaagat catcatcaaa acctgtggta ctactaagct 60
attgottgca atcccaccca tattgaagtt cgctgaaatg ctttccctta atgttaagtc 120
tgtgaattac accaggggaa gtttcatttt cccagtgct cagccatata cccatcgcaa 180
ctttttotgag gaagttgcta ttcttgatgg ctactttggc aaacttggtg caggaagcaa 240
tgottat 247

<210> 2518
<211> 336
<212> nucleic acid
<213> Glycine max
<400> 2518

gtgcangca cgcgtacgta agctcggaat tcggctcgag gccatatccc catcgcaact 60
tttctgagga agttgctatt cttgatggct actttggcaa acttggtgca ggaagcaatg 120
cttatatattt ggggtggccaa gacaaagcac agaactggca tgtctactct gcttctgcag 180

attctgtaac tcaatgtgac aatgtttaca ctcttgagat gtgcatgact ggcttgata 240
gagagaaagc acaggttttc tacaaagaac aatctgcttc agctgccatg atgactgtta 300
attccggcat tagaaaaatt cttccagatt ccgaga 336

<210> 2519
<211> 306
<212> nucleic acid
<213> Glycine max
<400> 2519

agtctatgc acgcgtacgt aagctcggaa ttccggctcga gatcaaaacc tgtggtacta 60
ctaagctatt gcttgcaatc ccacccatat tgaagtctgc tgaaatgctt tcccttaatg 120
ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgtcag ccatatcccc 180
atcgcaactt ttctgaggaa gttgctattc ttgatggcta ctttggcaaa cttggtgcag 240
gaagcaatgc ttatatatttg ggtggccaag acaaagcaca gaactggcat gtctactctg 300
cttctg 306

<210> 2520
<211> 247
<212> nucleic acid
<213> Glycine max
<400> 2520

catgtcgact cctatgttct gtctgagtc agtctctttg tttatgccta caagatcatc 60
atcaaaacct gtggtactac aaagctactg cttgcaatcc caccatatt gaaatttgct 120
gaaatgtttc cctcaatggt agatctgtga attacaccag gggaagtttc atctttcccg 180
gtgtcagcc ctatcccat cgcaactttt ctgaggaagt tgctattctt gatggctact 240
ttggcaa 247

<210> 2521
<211> 282
<212> nucleic acid
<213> Glycine max
<400> 2521

gtcncatgca cgcccaacgt aagctcggaa ttccggctcga ggtttacact cttgagatgt 60

gcatgactgg cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag 120
 ctgccatgat gactgttaat tccggcatta gaaaaattct tccagattcc gagatttgtg 180
 actttgactt tgnaccatgt gggtattcaa tgaactctgt tgaaggcgct gctgtttcta 240
 cgattcatgt taccacagaa gatgggttca gttatgcaag ct 282

<210> 2522
 <211> 305
 <212> nucleic acid
 <213> Glycine max
 <400> 2522

gcatgcacgc gtacgtaagc tcggaattcg gctcgagnaa aacctgtggt actactaagc 60
 tattgcttgc aatcccaccc atattgaagt tcgctgaaat gctttccctt aatgttaagt 120
 ctgtgaatta caccagggga agtttcattt tccccagtcg tcagccatat ccccatcgca 180
 acttttntga ggaagttgct attcttgatg gctactttgg caaacttggt gcaggaagca 240
 atgcttatat tttgggtggc caagacaaag cacagaactg gcatgtctac tctgcttctg 300
 cagat 305

<210> 2523
 <211> 287
 <212> nucleic acid
 <213> Glycine max
 <400> 2523

cgatatgcacg cgtacgtaag ctcggaattc ggctcgagct cttctgaagc atggccatgg 60
 cggtttctgc aattggtttt gaaggtttcg agaagaggct ggaaatatcc tttttccagc 120
 cgggactttt tgctgaccct gagggaatgg gttaagagc tcttgcaaag tcccagttgg 180
 atgagatact tacaccggct gcttgacca ttgtttcatc tctcagaaat gatcatgtcg 240
 actcctatgt tctgtctgag tccagtctct tgtttatgcc tacaaga 287

<210> 2524
 <211> 276
 <212> nucleic acid
 <213> Glycine max
 <400> 2524

tcgcangcac gcgtacgtna gctcggaatt cggctcgagg gcattagaaa aattcttcca 60
gattctgaga tttgtgactt tgactttgag ccatgtgggtt attcaatgaa ctctgttgaa 120
ggtgctgctg tttctacat tcatgttacc ccagaagatg gtttcagtta tgcaagcttt 180
gagactgttg gncatgactt caaagtgggtg aatctgaacg aaatgggttaa gaggggtattg 240
gcatgttttc tcccaactga gttcgtgttg cagttc 276

<210> 2525
<211> 302
<212> nucleic acid
<213> Glycine max
<400> 2525

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cccaactgag ttctctgttg 60
cagttcatgt ggatgggtgca agcnagttgt ttgatcagac gtgttttctg gatgttaagg 120
gatactgtcg cgaagagagg agccacgaag ggcttggaat ggggtggttct cttgtctacc 180
aaaaatttgc caagacttgt gactgtgggtt cacctagatc aactctgaag tgntggaaag 240
aggaagatga agaagagtag ttttcttaag tgtctttatt atgtccttgc gaaaataagt 300
cc 302

<210> 2526
<211> 274
<212> nucleic acid
<213> Glycine max
<400> 2526

cangcacgcg tacgtaagct cggaattcgg ctcgagaaaa cgataatgtc gactcctatg 60
ttctatctga gtccagcctc tttgtttatg cctacaagat catcatcaaa acctgtggta 120
ctactaagct attgcttgca atcccacca tattgaagtt cgctgaaatg ctttccctta 180
atgttaagtc tgtgaattac accaggggaa gtttcatttt cccagtgct cagccatatc 240
cccatcgcaa cttttctgag gaagttgcta ttct 274

<210> 2527
<211> 264
<212> nucleic acid
<213> Glycine max

<400> 2527

ctgcaattgg ttttgaaggt ttcgagaagn ngctggaaat atcctttttc cagccgggac 60
tttttgctga ccctgagggg atgggtttta gagctcttgc aaagtcccag ttggatgaga 120
tacttacacc ggctgcttgc accatangtt tcatctctca gaaatgatca tgtcgactcc 180
tatgttctgt ctgagtnacg tctctttgtt tatgcctaca agatcatcat caaaacctgt 240
ggtatacaaa gctactgttg cant 264

<210> 2528

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2528

ntgcacgcgt acgtaagctc ggaattcggc tcgaggctac tgcttgcaat cccacccata 60
ttgaaatttg ctgaaatgct ttccctcaat gttagatctg tgaattacac caggggaagt 120
ttcatctttc ccgggtgctca gccctatccc catcgcaact tttctgagga agttgctatt 180
cttgatggct actttggcaa gcttagtgca ggaagcaatg cttatatattt ggggtggccaa 240
gacaaatcac agaactggca tgtctactct gcttctgcag attctgtaa 289

<210> 2529

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2529

tcnnangcac gcgtacgtna gctcggaatt cggctcgagg cttgcaatcc cacccatatt 60
gaagttcgct gaaatgcttt cccttaatgt taagtctgtg aattacacca ggggaagttt 120
cattttcccc agtgctcagc catatcccca tcgcaacttt tctgaggaag ttgctattct 180
tgatggctac tttggcaaac ttggtgcagg aagcaatgct tatattttgg gtggccaaga 240
caaagcacag aactggcatg tctactctgc ttctgcagat tctgtaactc aatgtgacaa 300
tgtttacatc t 311

<210> 2530

<211> 308

<212> nucleic acid

<213> Glycine max

<400> 2530

tntangcac gcgtacgtna gctcggaatt cggctcgagc aggaagcaat gcttatatnt 60
tgggtggcca agacaaatca cagaactggc atgtctactc tgcttctgca gattctgtaa 120
ctocatgoga caatgtttac actctagaga tgtgcatgac tggcctggat agagagaaaag 180
cacaggTTTT ctacaaagaa caatctgctt cagctgccat tatgactgtt aattctggca 240
ttagaaaaat tcttcagat tctgagattt gtgactttga ctttgagcca tgtggttatt 300
caatgaac 308

<210> 2531

<211> 292

<212> nucleic acid

<213> Glycine max

<400> 2531

tgcangcac gcgtacgtaa gctcggaatt cggctcgagc aaagcacaga actggcatgt 60
ctactotgtt tctgcagatt ctgtaactca atgtgacaat gtttactc ttgagatgtg 120
catgactggc ctggatagag agaaagcaca ggTTTTctac aaagaacaat ctgcttcagc 180
tgccatgatg actgttaatt cgggcattag aaaaattctt ccagattccg agatttgtga 240
ctttgacttt gaaccatgtg gttattcaat gaactctgtt gaaggcgtg ct 292

<210> 2532

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 2532

cnaangcac gcgtacgtaag ctcggaattc ggctcgaggc agattctgta actcaatgtg 60
acaatgttta cactcttgag atgtgcatga ctggcctgga tagagagaaa gcacaggttt 120
tctacaaaga acaatctgct tcagctgcc tgaatgactgt taattccggc attagaaaaa 180
ttcttcacaga ttccgagatt tgtgactttg actttgaacc atgtggttat tcaatgaact 240
ctggtgaagg cgctgctgtt tctacgattc atgttactcc agaag 285

<210> 2533

<211> 326
 <212> nucleic acid
 <213> Glycine max

 <400> 2533

 gatgcagcgt acgt nagctc ggaattcggc tcgagaatcc tctctttgcg attgtgtttt 60
 gatctgtctc ctactgcgat agtttctcta ctgttacatg gccatggcgg tttccgcaat 120
 tggttttgaa ggtttctgaga aaaggttgga aatatccttt ttccagccgg gactttttgc 180
 tgacctgaa ggaaggggtc taagggctct taaaaatcc cagttgggtg agatctaaca 240
 ccagctgctt gcaccattgt ttcttcgctc aaaaaacgat aatgtcgact cctatgttct 300
 atctgagtcc agcctctttg tttatg 326

<210> 2534
 <211> 502
 <212> nucleic acid
 <213> Glycine max

 <400> 2534

 ccacgcgtcc ggatagttcg ttgcatgcaa ccatagtttt attaggattt tttcttcttt 60
 gntttcaatt aggttgtttt gntgctctcc ttcaaactcc atctttccaa atcctctctt 120
 tgcgattgng ttttgatctg ctctctactg cgatagnttc tctactgnta catggccatg 180
 gcggtttccg caattggttt tgaaggnttc gagaaaaggn tggaaatatc ctttttccaa 240
 ccgggacttt ttgctgacct tgaaagaagg ggtctaaang gctnttacao aatccaagtg 300
 ggtgagattc taacaccagc tgnttgnacc attggttctt ngctnaaaaa cgatnatgnc 360
 cacttctatg gtctatctna gttcangctt tttgggtatg cctaccaaga tcattattna 420
 aaactnnggg accacctaac tattgggttn aatcccccnt atttgaaatt gcttnaanng 480
 ctttccotta aggttaaact gg 502

<210> 2535
 <211> 291
 <212> nucleic acid
 <213> Glycine max

 <400> 2535

 nangtcgan gcacgcgtac gtaagctcgg aattcggctc gaggtactg cttgcaatcc 60

cacccatatt gaaatttgct gaaatgcttt ccctcaatgt tagatctgtg aattacacca 120
 ggggaagttt catctttccc ggtgctcagc cctatcncca tcgcaacttt tctgaggaag 180
 ttgctattct tgatggctac tttggcaagc ttagtgcagg aagcaatgct tatattttgg 240
 gtggccaaga caaatcacag aactggcatg tctactctgc ttctgcagat t 291

<210> 2536
 <211> 308
 <212> nucleic acid
 <213> Glycine max
 <400> 2536

gcangcagc gtacgtaagc tcgggaattc ggctcgagng ccaagacaaa gcacagaact 60
 gggcatgtct actctgcttc tgcagattct gtaactcaat gtgacaatgt ttacactctt 120
 gagatgtgca tgactggcct ggatagagag aaagcacagg tttttacaa agaacaatct 180
 gcttcagctg ccatgatgac tgttaattcc ggcattagaa aaattcttcc agattccgag 240
 atttgtgact ttgactttga accatgtggg tattcaatga actctgttga aggcgctgct 300
 gttttctac 308

<210> 2537
 <211> 308
 <212> nucleic acid
 <213> Glycine max
 <400> 2537

acgtcgcang cacgcgtacg taagctcgng aattcggctc gagggcaagc ttagtgcagg 60
 aagcaatgct tatattttgg gtggccaaga caaatcacag aactggcatg tctactctgc 120
 ttctgcagat tctgtaactc catgcgacaa tgtttactct ctagagatgt gcatgactgg 180
 cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag ctgccattat 240
 gactgttaat tctggcatta gaaaaattct tccagattct gagatttgtg actttgactt 300
 tgagccat 308

<210> 2538
 <211> 281
 <212> nucleic acid
 <213> Glycine max

<400> 2538

gtgcgatgca cgcgtacgta agctcggaat tcggctcgag gtactacaaa gctactgctt 60

gcaatcccac ccatattgaa atttgctgaa atgctttccc tcaatgtag atctgtgaat 120

tacaccaggg gaagtttcat ctttcccggt gtcagccct atccccatcg caacttttct 180

gaggaagttg ctattcttga tggctacttt ggcaagetta gtgcaggaag caatgcttat 240

attttgggtg gccaaagaaa atcacagaac tggcatgtct a 281

<210> 2539

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2539

antgcgatgc acgcgtacgt nagctcgga ttcggctcga gattctggca ttagaaaaat 60

tcttccagat tctgagattt gtgactttga ctttgagcca tgtggttatt caatgaactc 120

tgttgaaggt gctgctgttt ctaccattca tgttacccca gaagatgggt tcagttatgc 180

aagctttgag actgttgggt atgacttcaa agtgggtgaat ctgaacgaaa tggttaagag 240

gggatgggca gttttcttcc caatgagttc tctgttgag ttcagtggga ggtgcaaca 299

<210> 2540

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2540

ntgnncgcgt acgt nagctc ggaattnggc tcgagctcga gccgttcgag aagaggctgg 60

aaatatcctt tttccagcng gnactttttg ctgaccctga ggncatgggt tnangagntc 120

ttgcaaagtc ccagttggat gagatannta nacncgctgc ttgcaccatt gtttcatctc 180

tcagaaatga tcatgncgan tcctatgtnc tgtctgagtc cagtntcttn gtntatgcct 240

acaagatcat catcaaaacc tgnngtacta caaagctact gctt 284

<210> 2541

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 2541

gcgtacgtaa gctcggaatt cggctcgagn ttaggttttt ttgttgctct ccttcaaact 60
ccatctttcc aaatcctctc ttgcgattg tgttttgatc tgcttctac tgcgatagtt 120
tctctactgt tacatggcca tggcggtttc cgcaattggt tttgaagggt tcgagaaaag 180
gttggaataa tcctttttcc agccgggact ttttgctgac cctgaaggaa ggggtctaag 240
ggctcttaca aaatcccagt tgggtgagat tctaacacca gctgcttgca ccatggt 297

<210> 2542

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2542

tcgnngcacg cgtaacgtaag ctcggaattc ggctcgaggg ccaagacaaa tcacagaact 60
ggcatgtcta ctctgcttct gcagattctg taactccatg cgacaatggt tacactctag 120
agatgtgcat gactggcctg gatagagaga aagcacaggt tttctacaaa gaacaatctg 180
cttcagctgc cattatgact gttaattctg gcattagaaa aattcttcca gattctgaga 240
tttgtgactt tgacttgagc catgtgggta ttcaatgaac tctgttgaag gtgctgct 298

<210> 2543

<211> 390

<212> nucleic acid

<213> Glycine max

<400> 2543

ttcttgatat ccatccagat tgatagttca ttgcatgcaa ccatagtttt attaggtttt 60
ttcttctttg ttttcaatta ggtttttgct gctctccttc aaactccgtc tttccgaatc 120
ctctctttgt gattgtgttc tggtctgctt cctaccgca tagtttctct tctgaagcat 180
ggccatggcg gtttctgcaa ttggttttga aggtttcgag aagaggctgg aaatatcctt 240
tttccagccg ggactttttg ctgaccctga gggaatgggt ttaagagctc ttgcaaagtc 300
ccagttggat gagatactta caccggtgc ttgcaccatt ggttcatctc tcagaaatga 360
tcatgtcgac ttctaaggtc tggtgaanc 390

<210> 2544

<211> 284
 <212> nucleic acid
 <213> Glycine max

 <400> 2544

 nncngcacgc nnacgtaagc tcggaattcg gctcgagggg aagtttcatt ttccccagtg 60
 ctcagccata tccccatcgc aacttttctg aggaagttgc tattcttgat ggctactttg 120
 gcaaacttgg tgcaggaagc aatgcttata ttttgggtgg ccaagacaaa gcacagaact 180
 ggcatgtcta ctctgcttct gcagattctg taactcaatg tgacaatgtt tacactcttg 240
 agatgtgcat gactggcctg gatagagaga aagcacaggt tttc 284

<210> 2545
 <211> 295
 <212> nucleic acid
 <213> Glycine max

 <400> 2545

 gtgcnnngca cgcgtacgta agctcggaat tcggctcgag ctcagccata tccccatcgc 60
 aacttttctg aggaagttgc tattcttgat ggctactttg gcaaacttgg tgcaggaagc 120
 aatgcttata ttttgggtgg ccaagacaaa gcacagaact ggcatgtnta ctctgcttct 180
 gcagattctg taactcaatg tgacaatgtt tacactcttg agatgtgcat gactggcctg 240
 gatagagaga aagcacaggt tttctacaaa gaacaatctg cttcagctgc catga 295

<210> 2546
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2546

 ngtcgcatgc acgcgtacgt aagctcgga ttcggctcga gtgctcagcc atatccccat 60
 cgcaactttt ctgaggaagt tgctattctt gatggctact ttggcaaact tgggtgcagga 120
 agcaatgctt atattttggg tggccaagac aaagcacaga actggcatgt ctactctgct 180
 tctgcagatt ctgtaactca atgtgacaat gtttacctc ttgagatgtg catgactggc 240
 ctggatagag agaaagcaca ggttttctac aaagaacaat ctgcttcagc tggccatgat 300
 gactgttaat 310

<210> 2547
 <211> 374
 <212> nucleic acid
 <213> Glycine max

<400> 2547

cacttaaaant cgatgcacgc gtacgtaagc tcggaattcg gctcgagctt gcaccattgt 60
 ttcatctctc agaaatgac atgtcgacnc ctatgttctg tctgagtcca gtctctttgt 120
 ttatggctta caagatcatc atcaaaacct gtggtactac aaagctactg cttgcaatcc 180
 caccatatt gaaatttgct gaaatgcttt ccctcaatgt taatctgtga attacaccag 240
 gggaagtttc atctttcccg gtgtcagcc ctatcccat cgcaactttt ctgaggaagt 300
 tgctattctt gatggctatt tggcaagctt agtgcaggaa gcaagcttat atttgggtgg 360
 ccagacaaat caca 374

<210> 2548
 <211> 343
 <212> nucleic acid
 <213> Glycine max

<400> 2548

nnntannccc canacgtcgc angcacgcnt acgtnacgtc ggaattcggc tcgagntctt 60
 ctttgttttc aatnaggttt ttttgttgct ctcttcaaa ctccatcttt ccaaatectc 120
 tctttgcgat tgtgttttga tctgttctt actgogatag tttctctact gttacatggc 180
 catggcggtt tccgcaattg gttttgaagg tttcgagaaa aggttggaaa tctcttttt 240
 ccagccggga ctttttgctg accctgaagg aaggggtcta agggctctta caaatccca 300
 gttgggtgag attctaacac cagctgcttg caccattggt tct 343

<210> 2549
 <211> 292
 <212> nucleic acid
 <213> Glycine max

<400> 2549

nnnnnggcng acgtcgcang cacgcgtacg taagctcgga attcggctcg agcttacacc 60
 ggctgcttgc accattgtnt catctctcag aaatgaacat gtcgactcct atgtgctgtc 120

tgagtnacgt ctctntgttt atgcctacaa gatcatcatc aaaacctgtg gtactanaaa 180
gctactgctt gcaatccan ccatattgan atntgctgna atgctttccc ncaatgnag 240
atctngaat tacaccaggg gaagtttctt cttncctggg gctcagccct at 292

<210> 2550
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2550

ngtcgcatgc acgcntacgt aagcncggga attcggctcg agcaaatcc cagttgggtg 60
agattctaac accagctgct tgcaccattg tntcttcgct caaaaacgat aatgtcgact 120
cctatgtncct atctgagtec agcctctttg tttatgccta caagatcatn atcaaaacct 180
gtggtactac taagctattg cttgcaatcc cacccatatt gaagttcgct gaaatgcttt 240
ccctaagtgt aagtctgtga attacaccag gggaagtttc atttccccgt gctcagccat 300

<210> 2551
<211> 291
<212> nucleic acid
<213> Glycine max
<400> 2551

nngtcgcatg cacgcgtacg taagctcgga attcggctcg aggttcatgt ggatgggtgca 60
agcaagtcgt ttgagcagac ctgctttctg gatgttaagg gatactgtcg tgaagagagg 120
agccacgaag ggcttggaat ggggtggttct gttgtctacc aaaaattcgg gaagactagt 180
gactgtgggt cacctagatc aactctaaag tgctggaatg aggaagatga ggaagagtag 240
tttctttaag tgtctttatt attctgtcct gtgaaaataa gtctgggtttt c 291

<210> 2552
<211> 294
<212> nucleic acid
<213> Glycine max
<400> 2552

acgtcgcatg cacgcgtacg taagctcgga attcggctcg agtccaaatc ctctctttgc 60
gattgtgttt tgatctgctt cctactggcg atagtttctc tactgtttaca tggccatggc 120

ggtttccgca attggttttg aagggtttcga gaaaagggtg gaaatatcct ttttccagcc 180
 gggacttttt gctgacctg aaggaagggg tctaagggtt cttacaaaat cccagttggg 240
 tgagattcta acaccagctg cttgcaccat tgtttcttcg ctcaaaaacg ataa 294

<210> 2553
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2553

nangnacgg tncgtnacgt cacgtatanc tcggcattcg gctcgagctt tgttttcaat 60
 taggtttttt tgttgctctc cttcaaactc catctttcca aatcctctct ttgcgattgt 120
 gttttgatct gcttctact gcgatatgtt ctctactgtt acatggccat ggcgggtttcc 180
 gcaattgggt ttgaagggtt cgagaaaagg ttggaaatat cttttttcca gccgggactt 240
 tttgctgacc ctgaaggaag ggggtctaagg gctcttaca aatcccagtt gggtgaga 298

<210> 2554
 <211> 274
 <212> nucleic acid
 <213> Glycine max
 <400> 2554

acgcgtacgt aagctcggaa ttcggctcga gggtgcaagc aagtcgtttg agcagacctg 60
 ctttctggat gttaagggtt actgtcgtga agagaggagc cacgaagggc ttggaatggg 120
 tggttctgtt gtctacaaa aattcgggaa gactagtac tgtggttcac ctagatcaac 180
 tctaaagtgc tggaatgagg aagatgagga agagtagttt ccttaagtgt cttattattc 240
 tgtccttggtg aaaataagtc tggttttcca gata 274

<210> 2555
 <211> 263
 <212> nucleic acid
 <213> Glycine max
 <400> 2555

ncgtcgcatt cacgcgtacg taagctcggg attcggctcg agggtgcaag caagtcgttt 60
 gagcagacct gctttctgga tgtaaggga tactgtcgtg aagagaggag ccacgaaggg 120

cttggaatgg gtggttctgt tgtctaccaa aaattcggga agactagtga ctgtggttca 180
 cctagatcaa ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tccttaagtg 240
 tctttattat tctgtccttg tga 263

<210> 2556
 <211> 275
 <212> nucleic acid
 <213> Glycine max

<400> 2556

cacgcgtacg taagctcgga attcggctcg aggggtgcaag caagtcgttt gagcagacct 60
 gctttctgga tgttaagggg tactgtcgtg aagagaggag ccacgaaggg cttggaatgg 120
 gtggttctgt tgtctaccaa aaattcggga agactagtga ctgtggttca cctagatcaa 180
 ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tcctaaagtgt cttattattc 240
 tgtcctgtga aaataagtct ggttttccag atacg 275

<210> 2557
 <211> 280
 <212> nucleic acid
 <213> Glycine max

<400> 2557

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gaagcaatgc ttatatatttg 60
 ggtggccaag acaaatacaca gaactggcat gtctactctg cttctgcaga ttctgtaact 120
 ccattgcgaca atgtttacac tctagagatg tgcattgactg gcttgatag agagaaagca 180
 caggttttct acaaagaaca atctgcttca gctgccatta tgactgttaa ttctggcatt 240
 agaaaaattc ttccagattc tgagatttgt gactttgact 280

<210> 2558
 <211> 311
 <212> nucleic acid
 <213> Glycine max

<400> 2558

gtcgcangcg tacgtaagct cggaattcgg ctcgagccca gttgggtgag attctaacac 60
 cagctgcttg caccattgtt tcttcgtcnaa naaacgataa tgcgactcc tatgttctat 120

ctgagtcacag cctctttgtt tatgcctaca agatcatcat caaacctgt ggtactacta 180
agctattgct tccaatccca cccatattga agttcgctga aatgcttncc ttaatgttaa 240
gtctgtgaat acaccagggg aagtttcatt tccccagtcg tcagccatat ccccatcgca 300
atctttctgan g 311

<210> 2559
<211> 292
<212> nucleic acid
<213> Glycine max
<400> 2559

ctnecatgac gtaagtnago tcggaattcg gctcgagggt tttttgttg ctctccttca 60
aactccatct ttcacaaatcc tctcttttgcn ttgtgttttg atctgcttcc tactgcgata 120
gtttctctac tgttacatgg ccattggcggg ttccgcaatt ggttttgaag gtttcgagaa 180
aagggttgaa atattctttt tccagccggg actttttgct gacctgaag gaaggggtct 240
aagggtcttt acaaaatccc agttgggtga gattctaaca ccagctgctt gc 292

<210> 2560
<211> 288
<212> nucleic acid
<213> Glycine max
<400> 2560

ggntatgcac ncnacgtgn gcnngagtt cggcnngngg ggggnttccg naatnggttt 60
tgaagggttc gagaaaagg tggaaatatt cttntccan nccgggactt tntgctgacc 120
ctggnaggaa ggggtctaca gggctcttac aaaatcccag ttgggtgaga ttctaacacc 180
agctgcttgc accatgtttc ttnttcana aacgatnntg tcgactccta tgttctatct 240
gagtnacgcc tctttgttna tgctacaag atcatcancn anacctgt 288

<210> 2561
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2561

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc ncanngggg gagattcnaa 60

naccagctgc ttgcacnant gtttcttcgc tcaaaaacna gcaatgtcga ctcttatgtt 120
ctatctgagt ccagcctctt tgtttatgcc tncaaganca tcatcaaaac ctgggtacta 180
ctaagctatt gcttccaatc ccacccatat tgaagttcgc tgaaatgctt tcccttaatg 240
ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgtcag ccatatcccc 300

<210> 2562
<211> 236
<212> nucleic acid
<213> Glycine max
<400> 2562

ncgcgtacgt aagctcggaa ttccggctcga gcagcattcg ttgcggcatt ttaatcgatt 60
tatccaagca ggactgaatg aactaatgga gtctaaacnc gggaaaaaga agtctagtag 120
tagtagtagt aaatcattgt tctacgaagc tcccctcgga tacagcattg aagacgttag 180
accaaacggt ggaatcaaga aattcagatc tgctgcttac tccaactgcg ctcgcn 236

<210> 2563
<211> 285
<212> nucleic acid
<213> Glycine max
<400> 2563

ncncgtacgt aagctcggaa ttnggctcga gctcgagncg gttcatgtgg ntngtgcaag 60
caatttgtnt gatcanacgt gnnttctgga tgtaagga tactgtcgcg aagagaggag 120
ccacgaaggg cttggaatgg gtggttctct tnnctacca aaaatttgcc aagacttggt 180
actgtgcgct cacctagatc aactctgaag tgctggaaag aggaagatga agaagagtag 240
ttttcttaag tgtctttatt atgtccttgc gaaaataagt ccggt 285

<210> 2564
<211> 286
<212> nucleic acid
<213> Glycine max
<400> 2564

acgtcgcgtg cagcgtacg taagctcgga attcggctcg agaacttttc tgaggaagtt 60
gctattcttg atggctactt tggcaagctt agtgcaggaa gcaatgctta tattttgggt 120

ggccaagaca aatcacagaa ctggcatgtc tactctgctt ctgcagattc tgtaactcca 180
 tgcgacaatg ttacactct agagatgtgc atgactggcc tggatagaga gaaagcacag 240
 gttttctaca aagaacaatc tgcttcagct gccattatga ctgttg 286

<210> 2565
 <211> 296
 <212> nucleic acid
 <213> Glycine max

<400> 2565

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ngttttgggtt gattcaaggc 60
 cttcacagca ttcgntgagg ctttttnatn gatntntcca agcaggactg natgaactaa 120
 tggagtctaa aggtgggaaa aagaagtcta gtagtagtag tagtanatca ttttctacga 180
 agtcccccgc ggntacagca ttgaagangt tagaccaaac ggtggaatca agannttcag 240
 atctgtgtgt tactccaact gcgctcgcaa accttcctgn tatccatcca gattga 296

<210> 2566
 <211> 492
 <212> nucleic acid
 <213> Glycine max

<400> 2566

ccacgcgtcc gcttcaaadc acacactctc ttcaatttct agggttttgc tattgctttg 60
 cctccgttcc ccngntctca caaaaacaac gccttttctc ttctccttcg tatctattct 120
 ttgcgttttg tttttggttg attgaaggca ttcacagcng taattcgttg ctgcatttta 180
 atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagggtggg aaaaagaagt 240
 ctagtagtag tagtagtaaa tcaatttttt acgaagctcc cctcggatac agcattgaag 300
 acgttagacc aaacgggtgga atcaagaaat tcagatctgc tgcttactct aactgttctc 360
 gcaaaccatc ctgatacaca tccggattga tagttcgttg catgcaacca tagttttatt 420
 aggatttttt cttctttgnt ttcaattagg tttttttggt gctctccttc aaactccatc 480
 tttccaaatc ct 492

<210> 2567
 <211> 298
 <212> nucleic acid

<213> Glycine max

<400> 2567

gtagcatgca cgcgtacgta agctcggaat tcggctcgag cgtatctatt ctttgogttt 60
 gggttttggg tgattgaagg cattcacagc gtaattcggt gctgcatttt aatcgattta 120
 tctaaccagg actgaatgac ctaatggagt ctaaagggtg gaaaaagaag tctagtagta 180
 gtagtagtaa atcaattttt tacgaagctc cctcgata cagcattgaa gacgttagac 240
 caaacgggtg aatcaagaaa ttcagatctg ctgcttactc taactgttct cgcaaacc 298

<210> 2568

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2568

gtcgngcac gcgtacgtaa gctcggaatt cggctcganc tcgagccgaa tcggctcgag 60
 aganaagggt ggaaanctcc tttannccag ccgggacttt anggctgaac ctgaaggaag 120
 aggtctaagg gctcttacia aatcccagtt gggtagatt ctaacaccag ctgcttgac 180
 cattgtttct tcgctcaaaa acgataatgt cgactcctat gttctatctg agtcagcct 240
 cttgtttatg cctacnagat catcatcana acctgtg 277

<210> 2569

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 2569

ngtcgttgtn cgcgtacgtn agctcggnat tcggctcgag tttgttttca attaggtttt 60
 tgtctgctct ctttcaaact ccgtctttcc gaatcctctc tttgtgattg tnttctgttc 120
 tgcttcttac cgcgatagtt tctcttctga agcatggcca tggcggtttc tgcaattggg 180
 tttgaagggt tcgagaagag gctggaaata tcctttttcc agccgggact ttttgctgac 240
 cctgagggna tgggtttttag agctcttgcn aagtcccagt tggntgagat acttacaccg 300
 gctgctt 307

<210> 2570

<211> 245
<212> nucleic acid
<213> Glycine max

<400> 2570

ggtttttggct gctctccttc aaactccgtc tttccgaatc ctctctttgt gatttgtgttc 60
tgttctgctt cctaccgcga tagtttctct tctgaagcat ggccatggcg gtttctgcaa 120
ttggttttga aggtttcgag aagaggctgg aaatatcctt tttccagccg ggactttttg 180
ctgaccctga gggaaatgggt ttaagagctc ttgcaaagtc ccagttggat gagatactac 240
accgg 245

<210> 2571
<211> 326
<212> nucleic acid
<213> Glycine max

<400> 2571

nnncgcangc acgcgtacgt aagctcggaa ttcggctcga ggttttctcc caactgagtt 60
ctctgtttgca gttcatgtgg atggtgcaag caagttgttt gatcagacgt gttttctgga 120
tgtaagggga tactgtcgcg aagagaggag ccacgaaggg cttggaatgg gtggttctct 180
tgtctaccaa aatttgccaa gacttgtgac tgtggttcac ctagatcaac tctgaagtgc 240
tggaagagg nagatgcaga agagtagttt tcttaagtgt ctttattatg tcottgcgaa 300
aataagtcgg gttttccaga cagtga 326

<210> 2572
<211> 281
<212> nucleic acid
<213> Glycine max

<400> 2572

agttgcgcgn acncgtacgt aagctcggaa ttcggctcga gaaatcattg ttctangaag 60
ctccccctcn atanagcatn ggngacgtta gaccaanngg tggaatcaag aaattcagat 120
ctgntgctta ctccaactgc gctcgcaaan ctttctgat atccatcng attgatagtt 180
cattgcatgc aaccatagtt tnattaggtt ttntcttctt tgttttcaat taggtttttg 240
ctgctctcct tcaaactccg tctttccgaa tcctctcttt g 281

<210> 2573
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2573

 gtgcgcangca cgcgtacgta agctcggaat tcggctcgag gttttttctt ctttggtttc 60
 aattaggttt ttgctgctct ccttcaaact ccgtctttcc gaatcctctc tttgtgattg 120
 tgtttctgttc tgcttcttac cgcgatagtt tctcttctga agcatggcca tggcgggtttc 180
 tgcaattggt tttgaaggtt tcgagaagag gctggaaata tcctttttcc agccgggact 240
 ttttgctgac cctgagggaa tgggtttaag agctcttgca aagtcccagt tggatgag 298

<210> 2574
 <211> 450
 <212> nucleic acid
 <213> Glycine max

 <400> 2574

 aagggggatt ctgtangngn natngganng gacaacatan aagcgtatg acgtcgcatg 60
 cacgcgtacg taagctcgga attcggctcg agtttatgcc tacaagatca tcatcaaaac 120
 ctgtggtacn nntaagctat tgcttgcaat cccaccata ttgaagttcg ctgaaatgct 180
 ttcccttaat gttaagtctg tgaattacac caggggaagt ntcattttcc ccagtgtc 240
 nccatatcnn catcgcaagn tttntgagga agattnnant gttnttngtt antntnnonn 300
 ncttttttna tttacttaac ttatnatttg nncnttttat ntttaagcat natnactnna 360
 nntttttnag gnggggtgtn ttntttnttn ntctttnttn tttttttnnn attcanttta 420
 ttngttntn tntntntnn ntntctntt 450

<210> 2575
 <211> 218
 <212> nucleic acid
 <213> Glycine max

 <400> 2575

 annttacgnt tgaggtcang cacgcgtacg taagctcgga attcggctcg agnttttgaa 60
 ggtttcgaga aaaggttgga aatatccttt ttccagccgg gactttttgc tgaccctgaa 120

ggaaggggtc taagggctct tacaaaatcc cagttgggtg agatnctaac accanctgct 180
 tgnancattg tntcttcgct caaaaacgat aatgtcgn 218

<210> 2576
 <211> 428
 <212> nucleic acid
 <213> Glycine max
 <400> 2576

ccacnctcc gcaagttggt tgatcagacg tgttttctgn atgttaaggg atactgtcgc 60
 gaagnggggg gccactaagg gcttggaatg ggtgggttctc tnagtctacc aaaaatttgc 120
 caagacttgt gactgngggt cacctanato aactctgaag tgctggaaag aggaagatga 180
 anaanagtan ttttcttaag tgtctttatt atgtccttgc naaaataagt ccngttttnc 240
 agacagngat tgtttntctt tgggtgtttt tnccttttat gttagaccat tgtaggggca 300
 gtttggaacct tttattgntc tactattacc atttgaacat cgnatggatt ttaataaaan 360
 ananataata tnanngaaat ttatttctta ttattanctt ttnatntat ttnantttta 420
 naattctn 428

<210> 2577
 <211> 312
 <212> nucleic acid
 <213> Glycine max
 <400> 2577

togcatgcac gogtacgtaa gctcggaatt cggctcgagn taatgtcgac tcctatgttc 60
 tatctgagtc cagcctctnt gtttatgcct acaagatcat catcaaaacc tgtggtacta 120
 ctaagctatn gcttgcaatc ccacccatat tgnagtctgc tgaaatgctt cccttaatgt 180
 naagtctgtg aattacacca ggggaagnnt cattttcccc agtgcncagc canatncna 240
 togcaanttt tnngaggnag tcnccattcc tggatngcct actttggcaa acttggtgog 300
 gangcaatgc tt 312

<210> 2578
 <211> 261
 <212> nucleic acid
 <213> Glycine max

<400> 2578

gtcgcacatgca gcgtacgtaa gctcgggaatt cggctcgagc agccatatcc ccacgcgaac 60

ttttctgagg aagttgctat tcttgatggc tactttggca aacttggtgc aggaagcaat 120

gcttatatatt tgggtggcca agacaaagca canaactggc atgtctactc tggtttctgc 180

agattctgaa ctcaatgtgc caatgnttac actcntgagn tgnngcatga ctggctggat 240

agagagaang cncaggtttt c 261

<210> 2579

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 2579

gtcgcacatgca gcgtacgta agctcgggaat tcggctcgag taaagggtggg aaaaagaagt 60

ctagtagtag tagtagtaaa tcaatttttt acgaagctcc cctcggntac agcattgaag 120

acgttagacc aaacgggtgga atcaagaaat tcagatctgc tgcttactct aactgttctc 180

gcaaaccatc ctgatacaca tccggattga tagttcgttg catgcaacca tagttttatt 240

aggatttttc tcttgttttc aattagggtt tttgtgtgc 279

<210> 2580

<211> 234

<212> nucleic acid

<213> Glycine max

<400> 2580

tcgangcacg cgtacgtaag ctcggaattc ggctcgagaa gcaatgctta tattttgggt 60

ggccaagaca aagcacagaa ctggcatgtc tactctgctt ctgcagattc tgtaactcan 120

tgtgacaatg ttacactct tgagatgtgc atnactggcc tggatagaga gaaagcacag 180

gttttcnaca aagaacaatc tgcttcagct gccatgatga ctgnnaannc cggc 234

<210> 2581

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2581

gcacgcacgc gtacgtaagc tcggaattcg gctcgagntt cttctttgtt ttcaattagg 60
 tttttttgtt gctctccttc aaactccatc tttccaaatc ctctctttgc gattgtgttt 120
 tgatctgctt cctactgcga tagtttctct actgttacat ggccatggcg gtttccgcaa 180
 ttggttttga aggtttcgag aaaaggttgg aaatatcctt ttccagccgg gactttttgc 240
 tgaccctgaa ggaaggggtc taagggctct taaaaaatcc cagttgggtg agatctaaca 300
 ccagcn 306

<210> 2582
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 2582
 agtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ngccttcaca gcattcggtg 60
 cggcatttta atcgatttat ccaagcagga ctgaatgaac taatggagtc taaaggtggg 120
 aaaaagaagt ctagtagtag tagtagtaaa tcattgttct acgaagctcc cctcggtaca 180
 gcattgaaga cgttagacca aacgggtggaa tcaagaanct tcagatctgc tgcttactcc 240
 aactgcgctc gcaaaccctc ctgatatcca tccggttgat agttncattg catgcnacca 300

<210> 2583
 <211> 292
 <212> nucleic acid
 <213> Glycine max

<400> 2583
 cgtcgcgtgc ncgcgtacgt aagctcggaa ttcggctcga ggttttattn ggattttntc 60
 ttctttgtnt tcaattaggt tttttgttg ctctccttac aaactaccat actttccaaa 120
 tcctctcttt gcgattgtgt ttgatctgc ttctactgc ganagtatcn ctactgttac 180
 atggccatgg cggtnnccgc aattggtttt gaaggtttcg agaaaagggt ggaaatatcc 240
 ttttnccagc cgggactttt tgctgaccct gaaggaaggg gtctaagggc tc 292

<210> 2584
 <211> 153
 <212> nucleic acid
 <213> Glycine max

<400> 2584

catgcacgcg tacgtaagct cggaattcgg ctcgagctca gaaatgatca tgtcgactcc 60
tatgtttctgt ctgagtcag tctctttgtt tatgcctaca agatcatcat caaaacctgt 120
ggtactacaa agctactgct tgcaatccca ccg 153

<210> 2585

<211> 474

<212> nucleic acid

<213> Glycine max

<400> 2585

tnnaactnta cgcgcccagg taccggtcaa agaattcccg ggtcgacca cgcgtcngta 60
cggctgcgag aagacgacag aagggtacgg ctgcgagaag acgacagaag gggacacgca 120
actattttctg actacgtttt gctctacgcc tctccctctc tctcaaaaat cgttctcttc 180
gatttttaggg ttttgttttg ctgctgcctc cgttcccccc ttctcataaa caacgcggtt 240
tctcttctgc ttcgtatcta ttctttgctt ttggttttgg ttgattcaag gccttcacag 300
cattcggtgc ggcatttttaa tcgatttatc caagcaggac tgaatgaact aatggagtct 360
aaaggtggga aaaagaagtc tagtagtagt agtagtaaat cattgttcta cgaagctccc 420
ctcgataca gcattgaaga cgtagacca aacggtggna tcaaagaaat tcaa 474

<210> 2586

<211> 80

<212> nucleic acid

<213> Glycine max

<400> 2586

cagccctatc cccatcgcaa cttttctgag gaagttgcta ttcttgatgg ctacttnggc 60
aagcattgct nggnagnggg 80

<210> 2587

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2587

tcgcatgcan ncgtagnaat aagctcnana attcggctcg agggcntgga atgggtggtt 60

ctctngtnta ncaaaaacct gccaaagactt gtgactgtgg ttcacctaga tcaactctga 120
 agtgctggaa agaggaagat gaagaagagt agttttctta agtgtcttta ttatgtcctg 180
 cgaaaataag tccggttttc cagacagtga ttgtttttct ttggtgtttt ttccctttta 240
 tgtttagacca ttgttagggc gtttggacct tttattgttc tactattacc attgaacatc 300
 gaa 303

<210> 2588
 <211> 267
 <212> nucleic acid
 <213> Glycine max
 <400> 2588

ctgcacgcgt acgtaagctc ggaattcggc tcgagggaaa caagtcgttt gagcagacct 60
 gctttctgga tgtaagggga tactgtncgt gaagagagga gccacgaagg gcttggaatg 120
 ggtgggttctg ttgtctacca aaaattcggg aagactantg actgtgggtca cctagatcaa 180
 ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tccttaagtg tctttattat 240
 tctgtccttg tgaaaataag tctgggtt 267

<210> 2589
 <211> 225
 <212> nucleic acid
 <213> Glycine max
 <400> 2589

ttcaatttct agggttttgc tatacgcttt gcctcgttc cccctttctc acaaaaacaa 60
 cgccttttct cttctccttc gtatctatc tttgcntttn gtttttggtt gattgaacgg 120
 cattcacagc gtaattgggtg ctgcattttn atcgatttat ctaancagga ctgantgacc 180
 taatggagtc taaaggtggg aanagaagt ctagtagtag tagta 225

<210> 2590
 <211> 469
 <212> nucleic acid
 <213> Glycine max
 <400> 2590

tntaactctc cngcgcacag gtanacggta tcagagtccc cggcctcgac ccaagcgtca 60

agcccacgcg tccgtacggc tgcgagaaga cgacagaagg ggacacgcaa ctattttctga 120
ctacgttttg ctctacgcct ctccctctct ctcaaaaatc gttctctgnc gatttttaggg 180
tttctgtttg ctgctgcctc cgttccccc ttctcataaa caacgcgttt tctcttctgc 240
ttcgtatcta ttctttgctt ttgggttttg ttgattcaag gccttcacag cattcgttgc 300
ggcattttta tcgattttat caagcaggac tgaatgaact aatggagtct aaagggtggga 360
aaaagaagtc tagtagtagt agtagtaa atcattgttcta cgaagctccc ctcgataca 420
gcattnaaag aacttngac caaacggtg aatcaaggaa attcagatc 469

<210> 2591
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 2591
cacatgcacg cgtacgtaag ctcggaattc ggctcgaggt tttattagga ttttntcttc 60
tttgttttca attaggtttt ttgtgtgctc tccttcaaac tccatctttc caaatcctct 120
ctttgcgatt gtgttttgat ctgcttcta ctgcgatagt ttctctactg ttacatgcc 180
tggcggtttc cgcaattggt ttggaagggt tcgagaaaag gttggaaata tccttttcca 240
gccgggactt ttgtctgacc ctgaaggaag gggcttaagg gctctacaaa acccagtt 298

<210> 2592
<211> 275
<212> nucleic acid
<213> Glycine max

<400> 2592
anacactctc ttgcaatttc tagggttttg gtattgcttt gcctccgttc cccctttctc 60
ncaaaaaanaa cgnnttttct cttctccttc gtatctattc ttgcntntc gctttnggnt 120
gantnaaggc attcanagcg taattngttn ctgcattttn atcgatttct ctaancagg 180
ctgantganc taatggagtc taaagggtgg anaaagaagt ntagtagtan tagtagtaaa 240
tgcnttttnt acgnagctcc cctcgatac agcat 275

<210> 2593
<211> 269
<212> nucleic acid

<213> Glycine max

<400> 2593

cgcgnnocgtg ngctcgggaat tcggctcgag cgcaaaccat cctgatacac atccggattg 60
atagtctcgtt gcatgcaacc atagttttat taggattttt tcttctttgt tttcaattag 120
gtttttttgt tgctctcctt caaactccat ctttccaaat cctctctttg cgattgtgtt 180
ttgatctgct tctactgcg atagtntctc tactgttaca tggccatggc ggtttccgca 240
attggttttg aaggtttcga gaaaagggtt 269

<210> 2594

<211> 155

<212> nucleic acid

<213> Glycine max

<400> 2594

acgncgcacc nacgcgtacg taagctcgga attcggtctg aggtgggtact acaaagctac 60
tgcttgcaat cccacccata ttgaaatttg ctgaaatgct ttccctcaat gttagatctg 120
tgaattacac caggggaagt ttcattcttc ccggn 155

<210> 2595

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2595

naaagtcgca ngcacgcgta cgtaagctcg gaattcggct cgagttcaaa tcacacactc 60
tcttcaattt ctagggtttt gctattgctt tgctccggtt ccccttttct cacaaaaaca 120
acgccttttc tcttctcctt cgtatctatt ctttgcgttt ngntttnggt tgattgangg 180
cattcacagc gtaattngtt gctgcatttt aatcgattta tctaaccagg actgaatgac 240
ctaattggagt ctaaagggtg gaaaaagaag tctagtagta gtagtagtaa atcaattttt 300
n 301

<210> 2596

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2596

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ctctcttcaa tttctaggggt ttgctattg ctttgctcc gttccccott tctcacaaaa 120
acaacgcctt ttctcttctc ctctgtatct attctttgcg nttggttttt ggttgattga 180
aggcattcac agcgtaattt gttgctgcat ttnatcgat ttatctaacc aggactgaat 240
gacctaatgg agtctaaagg tgggaaaaag aagtctagta gtagtagtag taaatcaatt 300
tattacgaag c 311

<210> 2597

<211> 314

<212> nucleic acid

<213> Glycine max

<400> 2597

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tctcttcaat ttctaggggt ttgctattgc tttgcctccg ntcccccttt ctcacaaaaa 120
caacgccttt tctcttctcc ttctgtatct ttttttnt ttngtttng gttgattgaa 180
ggcattcaca gcgtaattng ttgctgcatt tnnatcgatt tatctaacca ggactgaatg 240
acctaatgga gtctaaagggt gggaaaaaga agtctagtag tagtagtagt aaatcaattt 300
tttacgaagc tccc 314

<210> 2598

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 2598

cgtacgtaag ctcggaattc ggctcgagat cacacactct cttcaatttc tagggttttg 60
ctattgcttt gcctccgttc cccctttctc acaaaaacaa cgctttttct cttctccttc 120
gtatctattc tttgcntttt gtttttggtt gattganggc attcacagcg taattngttg 180
ctgcatttnn atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagggtggg 240
aaaaagaagt ctagtagtag tagtagtaaa tcaatctttt acgaagctcc cctcggatac 300
agca 304

<210> 2599
 <211> 238
 <212> nucleic acid
 <213> Glycine max

 <400> 2599

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 gcctccgttc cccctttctc acaaaaacaa cgccttttct cttctccttc gtatctattc 120
 tttgcgtttg gtttttggtt gattganggc attcacagcg taattngtng ctgcatttna 180
 atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagnnggg aaaaagaa 238

<210> 2600
 <211> 274
 <212> nucleic acid
 <213> Glycine max

 <400> 2600

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 tctcgcaa ac catcctgata cacatccgga ttgatagttc gttgcatgca accatagttt 120
 tattaggatt ttttcttctt tgttttcaat taggtttttt tgttgctctc cttcaaactc 180
 catctttcca aatcctctct ttgcgattgt gttttgatct gcttcctact gcgatagttt 240
 ctctactgtt acatggccat ggcgggtttcc gcaa 274

<210> 2601
 <211> 257
 <212> nucleic acid
 <213> Glycine max

 <400> 2601

 caaganattc agatctgctg ctactctaa ctgttctcgc aaaccatcct gatacacatc 60
 cggattgata gttcgttgca tgcaaccata gttttattag gatttnttct tctttgtttt 120
 caattagggt tttttgttgc tctccttcaa actccatctt tccaaatcct ctctttgcga 180
 ttgtgttttg atctgcttcc tactgcgata gtttctctac tgttacatgg ccatggcggt 240
 ttccgcaatt ggntntg 257

<210> 2602

<211> 259
 <212> nucleic acid
 <213> Glycine max

 <400> 2602

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 tgtttngctg ctgcctccgt tcccccttc tcataaaca cgcgttttct cttctgcttc 120
 gtatctattc tttgcttttg gttttggttg attcaaggcc ttcacagcat tcgttgcggc 180
 attttaatcg atttatccaa gcaggactga atgaactaat ggagtctaaa ggtgggaaaa 240
 agaagtctag tagtagtag 259

<210> 2603
 <211> 246
 <212> nucleic acid
 <213> Glycine max

 <400> 2603

 cgcacctott tctgattgcc ttcaaatac acactctctt caattttctag ggttttgcta 60
 ttgctttgcc tccgttcccc ctttctcaca aaaacaacgc cttttctctt ctccttogta 120
 tctattcttt gcgtttggtt tttggttgat tganggcatt cacagcgtaa ttngttgctg 180
 catttnaatc gatttatcta accaggactg aatgacctaa tggagtctaa aggtgggata 240
 nagaag 246

<210> 2604
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2604

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 tccctctctc tcaaaaatcg ttctcttcga ttttagggtt ttgttttgct gctgcctccg 120
 ttccccctt ctcataaaca acgcgttttc tcttctgctt cgtatctatt ctttgctttt 180
 ggttttggtt gattcaaggc cttcacagca ttcgttgcgg cattttaatc gntttatcca 240
 agcaggactg aatgaactaa tggagtctaa aggtgggaaa aagangtcta gtagtagtat 300
 agtaaatcat 310

<210> 2605
 <211> 290
 <212> nucleic acid
 <213> Glycine max

 <400> 2605

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 tcaaaaatcg ttctcttcga tttnagggtt ttgttttgcg gctgcctccg ttccccctt 120
 nctcataaac aacgcgtttt ctcttctgct tcgtatctat tctttgcttt tggttttggt 180
 tgattcaagg ccttcacagc attcgttgcg gcattttaat cgatttatcc aagcaggact 240
 gaatgaacta atggagtcta aaggtgggaa aaagaagtct agtagtagta 290

<210> 2606
 <211> 333
 <212> nucleic acid
 <213> Glycine max

 <400> 2606

 nanatgnacg cgtacgtaag ctcggaattc ggctcgagct ttctgattgc cttcaaataca 60
 cacactctct tcaatttcta gggttttgct attgctttgc ctccgttccc cctttctcac 120
 aaaaacaacg ccttttctct tctccttcgt atctattctt tgcgtttggt tttcgggttga 180
 ntgaaggcat tcacagcgta attngttgct gcatttnnat cgatttatct aagcaggact 240
 gantgagcca atggngtcta aangtgggaa aaagaagtct agtngtagta gtagtaaata 300
 aattttttac gaagctcccc tcggatacag cat 333

<210> 2607
 <211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 2607

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 ttcaatttct agggttttgc tattgctttg cctccgttcc ccctttctca caaaaacaac 120
 gccctttctc ttctccttcg tatctattct ttgcgtttng ctttngggtg actganggca 180
 ttcacagcgt aattngttgc tgcattitna tcgatttatc taancaggac tgantgacct 240

aatggagtcc aaaggtggga aaaagaagtc tagtagtagt agtagtaa at caatTTTTTta 300
cgaagcnccc ctg 313

<210> 2608
<211> 286
<212> nucleic acid
<213> Glycine max
<400> 2608

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aaaaatcggt ctcttcgatt ttaggggttt gttttgctgc tgcctccggt ccccccttct 120
cataaacaac gcgttttctc ttctgcttcg tatctattct ttgcttttgg ttttggttga 180
ttcaaggcct tcacagcatt cgttgcgga ttttaatcga tttatccaag caggactgaa 240
tgaactaatg gagtctaaag gtgggaaaan naagtctagt agtagt 286

<210> 2609
<211> 311
<212> nucleic acid
<213> Glycine max
<400> 2609

ncgtcgang cacgcgtacg tnagctcggn attcggtcgc agnnttgctc tacgctctc 60
cctctctctc aaaaatcggt ctcttcgatt ttaggggtnt gttttgctgc tgcctccggt 120
cccccttct cataaacaac gcgttttctc ttctgcttcg tnttattctn gcgttttgg 180
gatnggttga ntnaaggcct ncanagcatt cgntgcgga ttctaatacga nttatccaan 240
caggntgaa tnaactantg gagtctaaag gnnggacaaa gaagtctagt ngtngtagna 300
gtaaancatt g 311

<210> 2610
<211> 306
<212> nucleic acid
<213> Glycine max
<400> 2610

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tacgcctctc cctctctctc aaaaatcggt ctcttacgat tttaggggtt ngttttgctg 120

ctgnetccgn tcccccttg ctcataaaca acgcgttttc nontctgctt cgtatctatt 180
 ctttgcntnn ggntttgggt gattcaaggc cttcacagca ttogtngcgg catttnaatc 240
 gatttatcca agcangactg aatgnactaa tggagtctaa aggtgggaaa aagaagtcta 300
 gtagtg 306

<210> 2611
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2611

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 aatcgttctc ttcgatttta gggtnntgtt ttgctgctgc ctccgttccc cctttnctca 120
 taaacaacgc gttttctctt ctgcttcgta tctattctac gcttttggtt ttgggttgatt 180
 caaggccttc acagcattcg ttgnggcatt tnaatcgatt tatccaagca ggactgaatg 240
 aactnatgga gnctaaaggn gggaaaaagc agtntagtag cngtagcagt nacncatgtn 300
 ntcagnagn cccngc 316

<210> 2612
 <211> 329
 <212> nucleic acid
 <213> Glycine max
 <400> 2612

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 accattgttt cttegtctcan aancnntnaa tgtcgantcc tatgttctat ntgngtccag 120
 cctgctttgt ttatgtcnta naagatcatc atcgcaacct gtggtntctac tangctattg 180
 cttgnaatcc cacccatatt gnngttcgct ganatgcttt ccttgaatgt taagtctgtg 240
 aattacacca agggaagttt cattttnnnc atgctcagnc atntccccgt cgcaactttt 300
 ctgaggaagn tgctattctt ggatggcta 329

<210> 2613
 <211> 274
 <212> nucleic acid
 <213> Glycine max

<400> 2613

agttctcatgc acgcntacgt nnagctcgga attcggctcg agtgcaacca tagttttatt 60

aggatttttt cttctttggt ttcaattagg tttttttggt gctctccttc aaactccatc 120

tttccaaatc ctctctttgc gattgtgttt tgatctgctt cctactgcga tagttttctct 180

actgttacat ggccatggcg gtttccgcaa ttggttttga angtttcgag aaaaggttgg 240

aaatatcctt tttccagccg ggactttttg ctga 274

<210> 2614

<211> 275

<212> nucleic acid

<213> Glycine max

<400> 2614

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acacactctc ttcaatttct aggnntttgc tattgctttg cctccgttcc ccttttctca 120

caaaaacaac gccttttctc ttctccttcg tatctattct ttgcgtttgg tttttggttg 180

attgaaggca ttcacagcgt aatngttgc tgcattttaa tcgatttate taaccaggac 240

tgaatgacct aatggagtct aaagntggga aaaag 275

<210> 2615

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2615

ncgcangcac gcgtacgtaa gctcggaatt cggctcgagn tgccttcaaa tcacacactc 60

tcttcaattt ctagggtttt gctattgctt tgccctcggt ccccttttct cacaaaaaca 120

acgccttttc tcttctcctt cgtatctatt ctttgcgctt ggcttaggnt gactganggc 180

attcacagcg taattcggtg ctgcatttta atcgatttat ctaaccaggnt ctgantganc 240

tantgggctc taaaggtggg aaaaagaagt ctaagtagta gtagagtaaa tcaantnncc 300

nc 302

<210> 2616

<211> 294

<212> nucleic acid

<213> Glycine max
 <400> 2616
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 tgectcogtt ccccccttct cataaacaac gcgttttctc ttctgcttcg tatctattct 120
 ttgcttttng ttttggttga ttcaaggcct tcacagcatt cgttgcgga ttttaatcga 180
 tttatccaag caggactgaa tgaactaatg gagtctaaag gtgggaaaaa gaagtctagt 240
 agtagtagta gtaaattcatt gttctacgaa gctccctcgc gatacagcat tgaa 294

<210> 2617
 <211> 320
 <212> nucleic acid
 <213> Glycine max
 <400> 2617
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 ttcaaataac acactctctt caatttctag ggttttgcta ttgctttgcc tccgttcccc 120
 ctttctcaca aaaacaacgc ctttntctt ctcttcgta tctattcttt gcnntngnt 180
 ntnggttgat tganggcatt cacagcgtaa ttngttgctg cattttaatc gatttatcta 240
 accaggactg aatgacctaa tggagtctaa anntggggaa aangaagtct agtagtagta 300
 gtagtaaata aatttntacg 320

<210> 2618
 <211> 260
 <212> nucleic acid
 <213> Glycine max
 <400> 2618
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 acaacgcggt ttctcttctg ctctgtatct attctttgct tttggttttg gttgattcaa 120
 ggccttcaca gcattcgttg cggcatttta atcgatttat ccaagcagga ctgaatgaac 180
 taatggagtc taaaggtggg aaaaagaagt ctagtagtag tagtagtaaa tcattgttct 240
 acgaagctcc cctcggatac 260

<210> 2619

<211> 285
 <212> nucleic acid
 <213> Glycine max

 <400> 2619

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 cnatctcaaa aatcgtgctc ttcgatttta gggttttggn ttgctgctnc ctccgttcnc 120
 cccttctcat aaacaacgnc gttttctctt ctgnttcgta tctattcttt gcttttggtt 180
 ttggttgatt caaggccttc acagcattcg gtgcggcatt ttaatcgatt tatccaagca 240
 ggactgaatg aactaatgga gtctaaaggt gggaaaaaga agtct 285

<210> 2620
 <211> 304
 <212> nucleic acid
 <213> Glycine max

 <400> 2620

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 gctgcctcgg tccccccctt ctcataaaca acgcgttttc tcttctgctt cgtatctatt 180
 ctttgctttt ggttttggtt gattcaaggc cttcacagca ttcgttgagg cattttaatc 240
 gatttatcca agcaggactg aatgaactaa tggantcnaa nggtgggaaa aagaagtcta 300
 gtag 304

<210> 2621
 <211> 405
 <212> nucleic acid
 <213> Glycine max

 <400> 2621

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 tgtgaggtag actcgtggaa gtttcatttt ccctggggca cagtcttttc ctcaccgcag 120
 tttttccgag gaggtttctg ttcttgacag ctatttcagc aaccttggtt ctggtagcaa 180
 agcatatggt atgggtgacc cttcaaagtc acagatttgg cacatctact ctgcaagtgc 240
 acagacaaaa ggatcatctg aagctgtcta tggcctagag atgtgcatga ccggtttaga 300

caaggaaagt gcttctgtgt ttttcaagga gaatacatct tcagcagctt caatgaccga 360
aaattctgga attangaaga ttcttccaca gtctgatata tctga 405

<210> 2622
<211> 299
<212> nucleic acid
<213> Glycine max
<400> 2622

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tcaagttggc tgatgctctt gacatagctg tgaaatctgt gaggtacact cgtggaagct 120
tcattttccc tggggcacag tcttttcttc accgcagttt ttccgaggag gtttctgttc 180
ttgacagcta tttcagcaac cttggttctg gtagcaaagc atatgttncg ggtgaccctt 240
caaagtcaca gatttggcac atctactctg caagtcaca gaccaaagga tcatctgaa 299

<210> 2623
<211> 200
<212> nucleic acid
<213> Glycine max
<400> 2623

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gtgaaatctg tgaggtacac tcgtggaagc ttcatTTTTCC ctggggcaca gtcttttctc 120
caccgcagtt ttccgagga ggtttctgtt cttgacagct atttcagcaa ccttggttnt 180
ggtagcaaag catatgttat 200

<210> 2624
<211> 328
<212> nucleic acid
<213> Glycine max
<400> 2624

gtnncatgca cgcgtacgta agctcggaat tcggctcgag ctgagaatgg gatggtgagt 60
agtgaggggtg aaaccagttt cacttggtgc atgaagtttg gtggctcctc tgtggcttct 120
gctgatagga tgaaagaggt ggctaccctt atattgagtt ttcccagga gaggcctatt 180
gttgttctct ctgctatggg aaaaacaaca aacaagcttt tgctggctgg agagaaagct 240

gtgagttgtg gtgttatcaa tgtatcaagt attgaggagc tttgctttat aaaagacctg 300
catctaagga ctgtggatca gcttggtg 328

<210> 2625
<211> 254
<212> nucleic acid
<213> Glycine max
<400> 2625

caaatgcgga catttttgaa gcaacttatc cggcagtcgc caagagatta catggtgatt 60
ggctctctga tcttgcaatt gcaattgta caggcttcct tggaaaggcc cagaaatcat 120
gtgcagtgc aacactgggt agagggggca gtgatttgac agctacagca attgggaaag 180
cactaggggt acctgagatc caggtatgga aggatgttga tgggtgccta acctgtgac 240
caaatatata ccca 254

<210> 2626
<211> 297
<212> nucleic acid
<213> Glycine max
<400> 2626

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gtatatcagt tgatgttgta gctacaagtg aagttagtat ttccttgaca ttggatccat 120
caaagctatg gagcagagaa ctaattcaac aggaacttga ctatgttgtc gaagaactgg 180
aaaaaattgc agtagtaaat ctcttaaaga ccagatccat aatctctctc attggaaatg 240
ttcagagatc atcactaata ttggagaagg cctttcatgt tcttcgaact cttgggg 297

<210> 2627
<211> 299
<212> nucleic acid
<213> Glycine max
<400> 2627

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agtattgagg agctttgctt tataaaagat ctgcatctaa ggnctgtgga tcagcttggt 120
gtggacggat ctgttattgc aaagcatcta gaagaattgg agcaacttct gaaggggata 180

gctatgatga aagaattgac taaaaggact caggactatt tagtctcctt ggagaatgca 240

tgctcgactag gatcttgctg gcatatctta acaaaatagg tgtcaaggcc gccatacga 299

<210> 2628

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2628

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aaggctcaat caanagatga gtcttataca gctgcattaa atgctgtttt ggagaagcac 120

agtgcgaactg cacatgacat acttgancgg agataatctt gctactttct tgtctaaatt 180

gcatcatgat attagtaacc ttaaggcgat gcttcgtgca atatacatag ctggtcatgc 240

aacagagtcc ttacagattt tgttggtggga catggagaat aggtct 286

<210> 2629

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2629

gtcgcgtacg tnagctcgga attcggctcg agcaggcctc aacctgaaa gaagaattca 60

acattgattt gcgtgtaatg ggcatacttg gttcanagtc aatgcttctt agtcatgttg 120

gcattgactt agctagatgg agagaacttc gagaggaaag aggagaagtg gctaattgtg 180

aaaaatttgt tcaacatgta catggaaatc attttatacc aaacacagca ttagtggact 240

gcacagctga ctctgccatt gctggctatt actatgactg gttgcgcaa 289

<210> 2630

<211> 168

<212> nucleic acid

<213> Glycine max

<400> 2630

angcacgcgt acgtnagctc ggaattcggc tcgagngcaa atggatggat acaagggatg 60

tccttatcgt aaatcctact ggttctaatc aagttgatcc tgactatttg gaatctgagc 120

aaagacttga aaaatggtac tctttgaatc catgtaaggt aatcattg 168

<210> 2631
 <211> 207
 <212> nucleic acid
 <213> Glycine max

 <400> 2631

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 ggctctgcat ccnngtcta tgagacctgc tanagaaagt gatattcctg ttaggggttaa 120
 aaattcctac aaccctaaag ctccaggtac tctcnttgc tnnngacgngg gatatggnga 180
 gggctttttt acctnccttt tttggag 207

<210> 2632
 <211> 293
 <212> nucleic acid
 <213> Glycine max

 <400> 2632

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 aacgagttac acgtgtgtca tgaagtttgg cggctcctct gttgccaatg cagaaaggat 120
 gagagagggt gccaacctta ttctgagctt cccggaagag aggcctataa ttgtntctct 180
 tgccatggga aagacaacta acatgctgtt gctggctgga gaaaaagctg taagctgtgg 240
 nataactang gnngatagtt tgacgnanng gttttnaaaa attggnatcc ggg 293

<210> 2633
 <211> 270
 <212> nucleic acid
 <213> Glycine max

 <400> 2633

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 tctgttcaca aatttgggtg aacctgtgtg ggaacctctc aaagaataaa aatgtttgct 120
 gacataattc ttaaggatga ttcggggaga aaattgggtg ttgtctctgc aatgtcaaag 180
 gtgaaaaata tgatgtatga cttatccac aaggctcaat cagcgatga gtcttatata 240
 gctgcattgg attctgttta ggagaagcac 270

<210> 2634

<211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 2634

 nncacnccga cgcacgcgta cgtaagctcg gaattcggct cgagcccgga aatcatgtgc 60
 agtgacaaca ctgggaagag ggggcagtga ttgacggct acaacaattg ggaaagcact 120
 agggttgcct ganatccagg tatggaagga tgttgatggt gtcctaacct gtgatccaaa 180
 tatataccca aaagcagaac ctgttcctta ttgacattt gatgaggctg cagaactagc 240
 ttactttggt gctcaggttc tacatccaca gtctatgaga cctgccagag aaagtgatat 300
 tcctgttagg gtt 313

<210> 2635
 <211> 322
 <212> nucleic acid
 <213> Glycine max

 <400> 2635

 gtgcangca tcgtacgtna gctcggaatt cggctcgagn taaaaggact caggactatt 60
 tagtntcctt tggagaatgc atgtnnacaa ggatctttgc nggnacatnc gnaanaaaat 120
 aggtgtcaag gctcgccaat atgatgcatt tganattggt ttataanca actgacgact 180
 tcacanatgc ggacattttg gnngccactt atccagctgt tgcanagagn ttgcatggtn 240
 antagctctc cgctcctgca attgcaattg cnacggctcc ttggaaaggc ccggaactca 300
 ngtgcantnc caacactggg ac 322

<210> 2636
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2636

 gtgcacatgca cgcgtacgta agctcggaat tcggctcgag ctacaagtga agtcagtgtt 60
 tccttgacat tggatccatc aaagctatgg agcagagagc taattcagca ggaacttgac 120
 catgtttgtag aagaactcga gaaaatcgct gtggatgaatc tcctgcagaa tagatccatc 180
 atctctctca ttggaaatgt tcagagatca tcactaatat tggagaagggt tctgtattat 240

gctcccttta tttaaattag ttcatatcct aactttcact tatgtataaa gatttactga 300
 atttattaca 310

<210> 2637
 <211> 438
 <212> nucleic acid
 <213> Glycine max
 <400> 2637

ccacgcgtcc gtcccaaata tggaagaata tcttggaag caggtcttag aggctgtttc 60
 attgattgaa gttagaagga acttcattga agcattatcc agtccgtttg gaaggccagt 120
 tgaagctgat gctgtctttt gcaggaagga aacttttctt gctgcatctg gtgttttcac 180
 gttcctgaag cagaagaaat accgtgtgac agaaatgttg ggatccaatt gcaagatatg 240
 ggacttgagt tccacattgg aagtaatgag attctaattg gggttttata agaccttggg 300
 ctctctgact acaatgatta gcttttgtgg tatgattcac tcaaagttct tatcaattgg 360
 tattagagct tttcattatg tctntcaact ncaaaacanc nacntgggtg aataccttca 420
 aatttgnact gttnaagt 438

<210> 2638
 <211> 329
 <212> nucleic acid
 <213> Glycine max
 <400> 2638

tgcangcacg cgtacgtaag ctcggaattc ggctcgagnt tactcctaac aagaaggcaa 60
 attcaggacc acttgatcag tatttgaagt taagagctct tcaaaggcaa tcctatacac 120
 attacttcta tgaagcaact gtcggagctg gtcttccaat tgtagcact ttacgtggcc 180
 tccttgaaac tggagacaaa atattacaaa tcgaaggcat ctttagtggg actttgagtt 240
 acatatttaa taactttaaa gatggccggg cttttagtga ggtagtttct gaagcaaagg 300
 aagcaggtta tactgagcca gatccaaga 329

<210> 2639
 <211> 256
 <212> nucleic acid
 <213> Glycine max

<400> 2639

cagtatttga agttaagagc tcttcaaagg caatcctata cacattactt ctatgaagca 60
actgttggag ctgggtcttcc aattgttagc actttacgtg gcctccttga aactggagac 120
aaaatactgc aaatcgaagg catctttagt gggactttga gttacatatt taataacttc 180
aaagacggcc gggcttttag tgaggtagtt tctgaagcaa aggaagcagg ttatactgag 240
ccagatccaa gagatg 256

<210> 2640

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 2640

gtcgnnntnn gggtagctga gctcggnatt cggctcgagg ggcagtatta acaagcattg 60
ttttgaaacg taatgtgacc atgttggata tagcaagcac tcgcatgctt ggtcngtatg 120
gtttccttgc taaggngttt tcaatctttg aagagttagg catatcagtt gatgtttag 180
ctacaagtga agtcagtgtt tccttgacac tggnnccatc aaagctatgg agcagagagc 240
taattcagca ggcaagtga cttgaccatg ttgtagaaga actcg 285

<210> 2641

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2641

cgcngcnttg tacgtnagct cggnatctcg ctcgagggca gtattaacaa gcattgtttt 60
gaaacgtaat gtgaccatgt tggatatagc aagcactcgc atgcttggtc agtatggttt 120
ccttgctaag gtgttttcaa tctttgaaga gttaggcata tcagttgatg ttgtagctac 180
aagtgaagtc agtgtttctt tgacactgga tccatcaaag ctatggagca gagagctaatt 240
tcagcaggca agtgaactga ccatgttgta gaagaactcg ag 282

<210> 2642

<211> 527

<212> nucleic acid

<213> Glycine max

ggagtggaca gaaatgttat tgaga

265

<210> 2645
<211> 307
<212> nucleic acid
<213> Glycine max

<400> 2645

cangcacgcg tacgttagct cgggattcgg ctgagcaag cactcctggt gttagtgcct 60
cccttttcaa tgcattggct aaggccaata taaatgtccg tgctatagcg caagggttgtt 120
ctgagtacaa tattactggt gttgttaagc gagaggattg tataaaggct ttacgagctg 180
tccattccan attttatctc tcaagaacca ccatagcaat gggcattatt ggacctggat 240
taattgggag cacactactt gaccagctaa gggatcaggc ctcaaccctg aaagaagaat 300
tcaacat 307

<210> 2646
<211> 327
<212> nucleic acid
<213> Glycine max

<400> 2646

nngnngcann cagcggtacg tnagctcgga attcggctcg anntccgtgc tatagcgcaa 60
ggntgttctg agtacaatat tactgttggt gttaancgag aggnttgtat aaaggcttta 120
cgagctgtcc attccagatt tnatctctca agaaccacca tagcaatggg cattattgga 180
cctggattaa ttgggagcac actacttgac cagctanagg atcaggcctc aacctgaaa 240
gaagaattca ncattgattt gcgtgtaatg ggcatacttg gttcaaagtc aatgctotta 300
gtgatgttgg cattgacttn nctagn 327

<210> 2647
<211> 317
<212> nucleic acid
<213> Glycine max

<400> 2647

ngtnngnnta cgtnagctcg gaattcggct cgagcggatc tgttatttca aagcatctag 60
aagaattgga gcaacttctg aaggggatag ctatgatgaa agaattgact aaaaggactc 120

aggactatTT agtctcctTT ggagaatgca tgtcgacaag gatctttgct gcataTctta 180
 ataaaatagg tgtcaaggct cgccaatatn atgcatttga gattggTTTT ataacaactg 240
 acgacttcac aaatgCGgac atTTtggaag cnaCTtatnc agctgttgca aagagattgc 300
 atggtgattg gctctcc 317

<210> 2648
 <211> 334
 <212> nucleic acid
 <213> Glycine max
 <400> 2648

tcgcangcac gcgtacgtaa gctcgggaaT tcggctcgag ctcaTTtcca gagcatctag 60
 aagaattgga gcaacttctg aaggggatag ctatgatgaa agaattgact aaaaggactc 120
 aggactatTT agtctcctTT ggagaatgca tgtcgacaag gatctttgct gcataTctta 180
 ataaaatagg tgtcaaggct cgccaatatg atgcatttga gattggTTTT atagcaactg 240
 acgacttcac aaatgCGgnc atTTtggaag caacttatcc agctgttgca aagagattgc 300
 atggtgattg gctctccgat cctgcaattg caat 334

<210> 2649
 <211> 286
 <212> nucleic acid
 <213> Glycine max
 <400> 2649

ngcacgcgta cgtaagctcg gaattcggct cngnGnatgt ccctccgCGa gancggcccc 60
 tccatcgccg tcgtgggCGt catcggcgCC gncggccagg agttcctact cgtcctctc 120
 cgaccgCGac ttccctacc gctcctcat atgctggctt ccaagcGctc cgtgGCCgc 180
 cgcacacct tcgaggacag ggactacGtc gtccaggagc tcacGCCgga gagcttcGac 240
 ggtgtcGcat cgcGctcttc agcGCCggcg gctccatcag caagca 286

<210> 2650
 <211> 280
 <212> nucleic acid
 <213> Glycine max
 <400> 2650

gtnnangcac gcgtacgtaa gctcgggaatt cggctcgagg acgctccggt gggcgccgca 60
tcacctttna gnnccagggga ctacgtcgtg nccgagctga cggcgggagan cttcgatggc 120
gtcganacgc cgatctncan cgcgcncggc nccattagc aagtacttcn gccccatcnn 180
cgtnatcgg ggaacgggtgn tcgncgacan cagatccgcn tntcggatgg acnanaatgt 240
cccattggta atncccgaat caaaccgtn nccatgcaaa 280

<210> 2651
<211> 323
<212> nucleic acid
<213> Glycine max

<400> 2651

ncacgegtnc gtanacgtnn gnattcngct cgannngcgn nnnngctgggg cgccgcatca 60
cctttgagtn acaggggacta cgtcgtggag gagctgacgg cggagagctt cgatggcgtc 120
gacatcgcg cgttcagcgc cggcgggctcc attagcaagt acttcggccc catcgccgtc 180
gatcggggaa cgggtggctgt cgacaacagc tccgcgtttc ggatggacga gaatgtccca 240
ttggtaatc ccgaagtga cccggaagca atgcaaaaca tcaaagccgg aatggaaagg 300
gcgcatcat gctaacccta att 323

<210> 2652
<211> 402
<212> nucleic acid
<213> Glycine max

<400> 2652

gggaataaaa cctgctttct atttttctca acctaaaatc ccatccacca accttatgat 60
gatgttttca ctctctgttt cgcgccacaa ccacctcttc tcgggccctc tcccggcccg 120
gccaagccc aagcccagct tttcctcttc caggatccga atgtccctcc aagaaaacgg 180
ccctccatc gccgtcgtgg gcgtcaccgg cgcgcgcggc caggagtctc tctcgtcct 240
ctcggaccgc gacttccct acagctccat caaaatgctc gcgtccaagc gctccgctga 300
gcgcgcgcatc acctttgagg acagggacta cgtcgtggag gagctgacgg cggagaactt 360
cgatggcgtc gacatcgcg tcttcancgc cggcgggctc aa 402

<210> 2653

<211> 482
 <212> nucleic acid
 <213> Glycine max

<400> 2653

gggnngggnn nnnnttaact ttccagggcc cggtcaggaa aacccgggtc gacccacgcg 60
 tongtacggc tgcgagaaga cgacagaagg gggggacatg gaattggaag acaagttgga 120
 gtgtggtggt ctgttttaaa atccaacact taatctctct cttcgagcc taaaatccca 180
 atggcttcac tctctgtttt gcgccacaac cacctcttct cgggccccct ccggccccgc 240
 cccaagccca cctcctctc ctctccagg atccgaatgt ccctccgga gaacggcccc 300
 tccatcgccg tcgtgggctt caccggcgcc gtgggacagg agttcctctc cgtcctctcc 360
 gaccggcact tcccctaccg ctccattcat atgctggctt ccaagcgctc cgtgggcccc 420
 cgcattacct tcgagacaaa ggactaagtc gtccaagaac tcangccggg anaacttcaa 480
 cg 482

<210> 2654
 <211> 327
 <212> nucleic acid
 <213> Glycine max

<400> 2654

cangcacgcg tacgtaagct cggaattcgg ctcgagctct cttcgagcc taaaatccca 60
 atggcttcac tctctgtttt gcgccacaac cacctcttct cgggccccct ccggccccgc 120
 cccaagccca cctcctctc ctctccagg atccgaatgt ccctccgga gaacggcccc 180
 tccatcgccg tcgtgggctt caccggcgcc gtgggacagg agttcctctc cgtcctctcc 240
 gaccggcact tcccctaccg ctccattcat atgctggctt ccaagcgctc cgtgggcccc 300
 cgcattacct tcgaggacag ggactac 327

<210> 2655
 <211> 312
 <212> nucleic acid
 <213> Glycine max

<400> 2655

gtngcccnca cgcgtacgta agctcggaat tcggctcgag gagctttgcc acaagaaggg 60

gactttgctc tgtattgatg gtacatttgc aacaccattg aaccagaagg cccttgccct 120
 tggcgctgat ctgattctgc actccttaac aaaatacatg ggtggacatc atgatgtaag 180
 ttggcatggt caagatttag ttgatggaa aaacaacaca aatatgttac attttcaggg 240
 ttaagagtta agcgtcaagg agcattttcc tctttcaggc ccttggtggt tgcataagtg 300
 gttcaactaa gg 312

<210> 2656
 <211> 542
 <212> nucleic acid
 <213> Glycine max
 <400> 2656

gnnnangtnn ntntcnaana ngntaanctt tnnaaacctt ccatttnggt aanncnccggg 60
 gccacgcgtc cgcncacgcg tncgcccacg cgtccggaga gaaagagagg gagtaagtgg 120
 gtinnaggang gaaaactaaa gaaacaaacc taacacaaca caaatccttg aaacgacgac 180
 ggaaatggcc gtttcgagct cgcacatgcg tttcaccttt gagtgccgct ccgatcccgga 240
 tttctcgcgc cccccgcgt ccttcgacaa cctccgcccgc cgaaaacttc gctcctccgc 300
 aggatccggc gcggcggtttc acggcatctc ctccctcatc ctccgcttcc tcccaacttc 360
 cagcgccagc taagcaccaa ggcgcgccgc aactgcagca acatcggcgt cgcgcaaadc 420
 gtgcgcgctt cgtggtcgaa caacagcnac aactnttcgg ccgncggggc ttncgcgcgcg 480
 accgggggnc cgggcacgga cccgggtacg ggcttttccc gtngtcgtaa cgncaacaa 540
 gg 542

<210> 2657
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2657

cgtcgcangc acgcgtacgt ncagctcgga attcggctcg agcggatctt gacatgggtg 60
 atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaacccca 120
 cccttacggg tgccaacata cctgaactgt gccacatggc acaccggaag ggagtgcagg 180
 tgggtggtgga caacacgttc gcgcccattg tgctttcgcc agcgcgtctt ggtgctgatg 240

ttgtcgttca cagtatctcc aagttcatca gcggtggggc cgatatcatt gcaggagcgg 300
tgtgcg 306

<210> 2658
<211> 307
<212> nucleic acid
<213> Glycine max
<400> 2658

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gacgatctt gacatggttg 60
atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaacccca 120
cccttacggg tgcgaaacata cctgaactgt gccacatggc acaccggaag ggagtgaagg 180
tggtgggtgga caacacgttc gcgcccattg tgctttcgcc agcgcgtctt ggtgctgatg 240
ttgtcgttca cagtatctcc aagttcatca gcggtggggc cgatatcatt gcaggagcgg 300
tgtgcg 307

<210> 2659
<211> 515
<212> nucleic acid
<213> Glycine max
<400> 2659

gnnnnngnaga gtttgnttgg ggggnaggga gnnanatttt nagaccacta tgacgtcgca 60
tgcaacgcgta cgtaagctcg ganttcggct cgagagacag aggntagaga angagaggga 120
gtaagtgggt aaaggaaaga naactaaaga aacaaaccta acacaacaca aatccttgaa 180
acgacgacgg aaatggccgt ttcgagctcg cacatgcgtt tcacctttga gtgccgntcc 240
gatcccgatt tctcgcccc cccgccgtcc ttcgacaacc tccgccgcg caacttccgc 300
tcttcgcag gatcccggcg cggcggttca cggcatctcc tctcactct ncgcttccct 360
cccaacttcc agcgccagct aagcaccaag gcgcgccgca actgcagcaa catcggcgtc 420
gcgcaaactcg tcgccgcttc gtggtcgaac aacagcgaca actctccggc cggcggggct 480
tcggcgccgg ccgcggcacc ggcacggacg ccgtt 515

<210> 2660
<211> 258
<212> nucleic acid

<213> Glycine max

<400> 2660

caaggacatc atgatgtcct tgggtggttgc ataagtgggt caattnnngt ggtttcgcaa 60
attcggactt tgcaccatgt nttgggtggt aactttaacc cgaatgctgc atacctatc 120
atcagaggca tgaaaacgct gcatctccgt gtacagcagc agaattcaac aggaatgagg 180
atggccaaac ttttagaggg acatcccaag gtgaagcggg tctactatcc aggcttgccg 240
agtcaccctg aacatgag 258

<210> 2661

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2661

ganattgtca tgcactctgc tacaaaatth attgctggac atagtgacat tatggctggt 60
gtgcttgctg tgaagggtga aaagtggga aaggaattgt atttcttgca aaatgcagag 120
ggttcaggct tagcaccatt tgactgttgg ctttgtttgc gaggaatcaa gacaatggcc 180
ctgcgaattg aaaaacaaca ggataatgca cagaagattg ctgagttcct tgctcccat 240
cctcgagtga agaaagtga ttatgctggc ttgctg 277

<210> 2662

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2662

gcatgcacnc gtacgtaagc tcggaattcg gctcgagntt gcctcccatc ctcgagtga 60
gaaagtgaat tatgctggct ngcctgggtca tcctggctgt gatttacact attctcaggc 120
aaagggtgca ggatctgtgc ttagcttctt gactgggtca ttggaacttt caaagcatat 180
tgttgaaact accaaatact tcagtataac cgtcagcttt gggagtgtga agtcccttat 240
tagcatgcca tgctttatgt cacatgcaag cnngncngct ggcagttcgt gaggccagag 300
tttaagaaga tctgtagtat at 322

<210> 2663

<211> 273
 <212> nucleic acid
 <213> Glycine max

 <400> 2663

 cgcangcacg cgtacgtaag ctcggaattc ggctcgagaa acaacaggat aatgcacaga 60
 agattgotga gttccttgcc tcccatcctc gagtgaagaa agtgaattat gctggcttgc 120
 ctggtcaccc tggctgtgat ttacactatt ctcaggcaaa gggtcgagga tctgtgctta 180
 gcttcttgac tggttcattg gaactttcaa agcatattgt tgaaactacc aaatacttca 240
 gtataaccgt cagctttggg agtgtgaagt ccc 273

<210> 2664
 <211> 289
 <212> nucleic acid
 <213> Glycine max

 <400> 2664

 gcaagcgtag gtaagctcgg aattcggctc gaggttcagg cttagcacca tttgactggt 60
 ggctttgttt gcgaggaatc aagacaatgg ccctgcgaat tgaaaaacaa caggataatg 120
 cacagaagat tgctgagttc cttgcctccc atcctcgagt gaagaaagtg aattatgctg 180
 gcttgccctgg tcctcctggg cgtgatttac actattctca ggcaaagggt gcaggatctg 240
 tgcttagctt cttgactggt tcattggaact tcaaagcata ttgttgaaa 289

<210> 2665
 <211> 499
 <212> nucleic acid
 <213> Glycine max

 <400> 2665

 aacttttacg ccnccangtg ccgggcaang gtnangagnt cccgggtcga cncacgcgtc 60
 cggcnagaag acgacagaag gggacgggtg tggcggacaa cacgttcgcg cccatgggtg 120
 tttcgccagc gcgtcttggt gctgatgttg tcgttcacag tatctccaag ttcacacagc 180
 gtggggccga tatcattgca ggagcgggtg gcggaccgcg aagactgggtg aacgcaatga 240
 tggatctgca acaagggtca ctaatgctgc tgggtccaac aatgaatgcg aaagtggcat 300
 tcgaactctc ggagagaata ccgcacctag ggctaagaat gaaggagcat agcaaccgcg 360

cactagagtt cgcaacgagg ctcaaaaggc taggaatgag ggtaatatatac ccgggcctgg 420
aggagcaccc acagcaccaa gcttctgaaa tcaatgcaca acaaggacta tggctacggc 480
gggctcatgt tcntngaca 499

<210> 2666
<211> 326
<212> nucleic acid
<213> Glycine max
<400> 2666

tgcgatgcac gcgtacgtaa gctcgggaatt cggtctgaga cgggaggtca agcccgcct 60
gaccaacatg gtctccgctg ctaagctcat tcgcacccag ctccgcagcg ccaagtgagc 120
accttttttt gccttttttcg ttcccgagga gggcgctcgtc gatgcccaatt tgtntccaat 180
aaacaggggt cccccctgt gcccgccgtt ctgttgtgct ccgtctgtgg ttaggttact 240
agttttcttg atctcgcccc cacgcgggtac ctgttttata tctgtttggg ggtttctgag 300
gcaagttgcc cgtgtattgt atcgta 326

<210> 2667
<211> 308
<212> nucleic acid
<213> Glycine max
<400> 2667

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gggacactac tcagattcac 60
actcacatgt gctactccaa cttcaatgac atcattcact caatcataga catggatgct 120
gatgtgatca ccattgagaa ctccagatca gatgagaagt tactttcggg cttccgtgag 180
ggagtgaat atggtgccgg cattggtcct ggcgtttatg atattcactc acccaggatt 240
cctcccacag aagaaattgc tgacaggatc aacaagatgc ttgcagttct tgaaagcagc 300
attctctg 308

<210> 2668
<211> 313
<212> nucleic acid
<213> Glycine max
<400> 2668

nagtcgcacg cacgcttacg taagctcnga attcggetcn nnnngnccant gancttgctg 60
 cttctctcan taccttggca gggctcttgag ggcattgggtg ggcaaagata agcttggttg 120
 tgtccacctn ctctccctt cttcacactg gctgtggntc tagttaacga naccnagttg 180
 gatgatgaga tcaagtcatg gctagctttg gctgccccaa aaattgttga ngttaacgca 240
 tnggctaaag cngtgtcngg ccacaaggat gaggcctnct tctctgntan tgcngctgct 300
 ctggettcaa gan 313

<210> 2669
 <211> 290
 <212> nucleic acid
 <213> Glycine max

Sequence 1

<400> 2669
 cgcangcacg cgtacgtaag ctcggaattc ggctcgaggg ggatgtactt gatcccannc 60
 accagccatc tgctcgacgc caccgccanc ctcggtgccg tcccccccag gtacggcttg 120
 accggggggg agattggatt cgnacactac ttctccatgg ccagaggtaa tgctaccgtg 180
 cntgctangg agatgaccaa gtggttcgac accaactacc actttattgt cctgaattg 240
 ggccctgatg tgaattcacc tatgcttctc acaaggctgt tgatgaatac 290

<210> 2670
 <211> 289
 <212> nucleic acid
 <213> Glycine max

<400> 2670
 cncgcacgcg tacgtaagcn tcgcgaattc ggctncgagt tgctcccnca gnagcganga 60
 agangccaca gagnnctagt ctccctanct ctacacccgc aaganaaana tggcatctca 120
 catacgttgg atacccccgt atgggttccc aagagagnnt ncaanttcgc tctcgagtcn 180
 ttctgggatn gcaanancag cgccgangct ttgcagangg tgtcntctga tctcagggca 240
 nccatctggn ancagttggc tgatgntggg ntcaagtaca tccccngca 289

<210> 2671
 <211> 303
 <212> nucleic acid
 <213> Glycine max

<400> 2671

nttgcgtnc a cgcgtacggn agctcggaat tcggctcgca gtgcacgtgt nattangnna 60
aatgagaaan aaaaaaatgg tcatencaca tcgnnggata cncctgtatg ggtgccaag 120
agagagcnca agntcgcttc tcgagtcttn cngggatggc aagagcagcg cngaggattt 180
cncagaaggt gtcttcngan ctcaaggcat ccancnggaa gcagatggct gttgctggga 240
tcaagtanan ccnancan acttcngcnc actatgatca ggntcacnac gncctgccac 300
nct 303

<210> 2672

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2672

agngacgcgt actaagtcgg aattcggctc gagaagataa gcttggttg tccacctct 60
cctcccttct tcacactgct gtnganccca gttaacgaga ccaagttgga tgatgagatc 120
aagtcattggg ctagcttttg nctggcccaa naaaattgtn ngaagttaaa cggcattggg 180
ctaaaaggca ttggtgctgg gaccancaa ggatggaggg cctttctttc tctgggtaaa 240
tggttggtg gctactgggc tttccaaggg aaagttcttt cctn 284

<210> 2673

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2673

cntgencgta nntnagctcg gaattcggct cgagctgagg ttgttggttg tgccaaggnt 60
gtaaatnatg gtcccgttc tgtgcgtgat ggctnctnaa nccaatgcca aatccatgca 120
ggngentgcc acgtcgaanc gtaccantaa caaggcgggt aaggaccgtc aagccagcgt 180
cactcctgag caacacngnc gcaagtctca gttccctgaa cgttatgcc agcagaagaa 240
gcacttgagc ctcttggttc cccaccacca ccattggttc cttccctcag accana 296

<210> 2674

<211> 269

<212> nucleic acid

<213> Glycine max

<400> 2674

gcgcntgcan tnanacgtng agntcggaat tcggctcgag cttccaatcc ttccaaccac 60
cactattggn ncnttcngtc agantnnann antgaggagg gtacgccgtg agttcaangc 120
taanaagntc tnnnnnggaa ntntatgnct aagtcaatta aggnnggaaat tcgcaaagtt 180
gttgaacttc aagaagagct tgatattgat gttcttggtc atggagancc agagngaaat 240
gntatggttg agtacttcgg tgagcaatt 269

<210> 2675

<211> 216

<212> nucleic acid

<213> Glycine max

<400> 2675

nngtcganac ntgcgtacgt aancncggaa ttcggctcga gggcaagacc agcgccgnng 60
attngcaggn ggtgtctnct aancncagga gcatccatch ggaagcagat ggcngatgct 120
gggatcaant acatccccag caacactttg ctactcacna naaccaggtn ctcgacgccn 180
ccgccaccct acggngccgn tgccacnnag gtangg 216

<210> 2676

<211> 263

<212> nucleic acid

<213> Glycine max

<400> 2676

anatgaccag aagacggtca ttggctttgg gcggctctgg ctgagagctt taccaagcgc 60
ccaatgaagg gaatgcttac cggaccagtt gagaatggta acaggaccgg taagcattct 120
caactggtcc ttgtttagaa atgaccaacc tagatctgag accacctacc agattgcttt 180
gtctatcaag gacgaagtgg aagacctga aaaggctggc atcactgtta tccaaattga 240
tgaagctgct ttgagagagg gtc 263

<210> 2677

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2677

cgcangcacg cgtacgtaag ctcggaattc ggctcgagtn gggngcgggn ggcgtgaagc 60

caccgatcat nnatgggtgat gtgagccgcc caaagccaat ganngtcttc tngngatnnc 120

tggntcagag ctttacnaag cgcccaatga agggaatgct taccggtcct gttaccannc 180

tcaactgggn nnntgttaga aatgaccanc ctagatctga gaccacctag nagantgctt 240

tgtctatcaa ggacgaatgg aaganncnhn ccaaggctgg catcantgtn a 291

<210> 2678

<211> 267

<212> nucleic acid

<213> Glycine max

<400> 2678

cctanggtgc cgntccacna nggnacngnt gncnccggcg gngagattng gtttgatacc 60

tactttctcca tggccanang taatgctacc gtgccagcta tggagatgac caagtgggttc 120

gacaccaact accactntat tgtccctgaa ttggggcctg atgtgaactt cacctatgct 180

tctcacaagg ctgttgatga atacaaggag gccaaaggcg ttgnagtgga taccgttccg 240

gtcctcgttg gccctgttac atacctg 267

<210> 2679

<211> 252

<212> nucleic acid

<213> Glycine max

<400> 2679

cagaagccnc agaagaagcc acagagaact agtctnctac tcnccaccng caagnaccnn 60

natggcatnn tcnctcggtt ggataccccc gtatgggtcc ncaannngag agtcaagtt 120

gtctcgagtc tttctgggat ggcaagagnn agcgccgagg gatttgcaga aggtgtcttc 180

tgatctccag ggcattccatc tggaagcaga tggatgatgct gggatcaagt acatccccag 240

caaactttct nt 252

<210> 2680

<211> 324

<212> nucleic acid

<213> Glycine max

<400> 2680

gtcgcangca cgcgtacgtn agctcggnat tcggctcgag cacgagacca agttggatga 60
tgagatcaag tcatggctag cttttgctgc ccaanaaatt gttgaagtta acgcattggc 120
taaagcattg tctggccaca aggatgaggc cttcttctct ggtaatgctg ctgctctggc 180
ttcaaggaag tcttctccaa agagttgacc aaacgagggc tgntccagaa agnctgctgc 240
tagcaattga agggttccag atcatngccg gncaacaatt ntccatgcc aactggatnc 300
tcaacaaaag aagnncaacc ttcc 324

<210> 2681

<211> 362

<212> nucleic acid

<213> Glycine max

<400> 2681

tnnaggtagt agctcggatt cggctcgagct ctccatggcc agaggtaatg ctaccgtgcc 60
tgotatggag atgaccaagt ggtncgacac naactaccac tttattgtgc ccngnattgg 120
gccttggatg nagaactttc acctatgnct ntcttcacaa gggctgtntg gatgcagata 180
ncaagggaag gtccaagggc cgatttgggg agtcgggata nccaatntcc ccggnaactc 240
cgtntggggc cnetgggtta ancatnactt tggtataggc tacctnncca aaggctcctg 300
gcacaaangg ggcaaatttc gganggtaaa attcccattg gggggcttcc tggccttgc 360
tc 362

<210> 2682

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2682

cgctcgnangc ccgcgtncgt cagctcggnn ttccgctcgn gctgnacctg tgcttgtggg 60
acctgtttct tncctgttgc tgtcannacc agctaagggt gttgagangt cattttccct 120
tctttcccta attgacaaga tcttctctgt ctacaggga gttgtggctg aactgaaggc 180
agctggtgct acttggatcc agtttgatga acctaccctt gtgaaggatc tcaatgcccc 240
ccagttacaa gcatttacc cagcctacgc agagtttagag tcaagtttnt ctggtttggn 300

tggttctgatt ganacatact t

321

<210> 2683
<211> 315
<212> nucleic acid
<213> Glycine max

<400> 2683

gnngcacgcg tacgtaagct cggaattcng ctcgagctgn cntcaacttg ntccctcaga 60

agcgnntann aatccacana gaactatgtc tcnctactnc ncacccgcaa gaaaaaaatg 120

gcntntcaac atcgttggat acccccgtat nggncccaat agaganctna attcgctctc 180

gagtctatct gggatgncaa nagtacgccg aggatattgca naangngtct actgatctca 240

gggcattcnt ctgganncag atggntnatg ctgggatcag tacatccnca gcaacatttc 300

tctnactntg accan 315

<210> 2684
<211> 174
<212> nucleic acid
<213> Glycine max

<400> 2684

gctgcaggca cgcgtacgtn agctcggaat tcggctccgn gcttcanctt gctccctcag 60

aagcgnagaa gaagcccana gagaanagnc tcctantctc acccgcaagn nnaanatggn 120

atctcacatc gttggntacn cccgtatggg tcccaagaga gagctnaagt tgnt 174

<210> 2685
<211> 303
<212> nucleic acid
<213> Glycine max

<400> 2685

aacngaaagn cgcangcacg cgtacgtaag ctcggaattc ggctcgagct tcaggatctc 60

cgaggaagag tatgtaaagt caattaagga ggaaattcgc aaagttgttg agcttcaaga 120

agagcttgat attgatgttc ttgttcattg agaaccagag gtccgctctc atttcataac 180

atgactaaat attagtcttt tgaattgaag atagcttctt tctttctgaa gagcaactac 240

tctttgcata ttttctttcc tattgttaga ttgcngattc caaaattgca gcacctccaa 300

ctn

303

<210> 2686
<211> 103
<212> nucleic acid
<213> Glycine max

<400> 2686

gntttntttt ttttttgggt tgttttcnnt tggttttttt tgtttttnnt gnttnttggt 60
tntttgtttt tttttgtttt ttntgtttt tttttttttt nnt 103

<210> 2687
<211> 523
<212> nucleic acid
<213> Glycine max

<400> 2687

gnnngnaggt tttgannggg ggggaggggn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnngnttgnt nccttanaag cnaagaagaa 120
nccncanana ancnggcttc taattttttt taaccaccag aaaaatgggn ttttaaantc 180
ggtnggntnc cccggattgg gnccaaggag agagctaaag ttcgttctcg agtctttctg 240
ggatggcaag gagcagcgcc gaggatttgc agaaagggtg ctgcttgatc tcaaggatcat 300
ccatctggna agcagatggc tgggtggctgg gatcaaagta ccttccccag caacactttt 360
ctcgttctat gaccaagctt gntcgacgcc accgncaccc tcgggtgccgt cccccccagg 420
tacggntgga ccggcggcga gattggattc gacacctact tctccatggn cagaggtaat 480
gctaccgtgc ctgctatgga gatgaccaat gggtcgacac caa 523

<210> 2688
<211> 570
<212> nucleic acid
<213> Glycine max

<400> 2688

agaggcctgc ttgnctttat canananann ncacngccat gccctccgaa cttccatttc 60
ngcccaggcg tactcccacg cntcctcagt ngccaacga ggctgttcan aanctgctg 120
ctgcattgaa ngnttnggan ggtntcctg caacaaatgt cagggtgccan actggattct 180

caacaaaaga anctcaacct tncaatnctg ncaaccacca ctattggatc cttccctcag 240
 actgtataac tgaggagggt acgcccctga attnaaagct aacaagatct ccgangaana 300
 ngtatgttna tgtcanttta aggaggaaat tttcaaantt tgtttgagct tnnataaaag 360
 cttnatattn atnttcttgt catnnagaan nncagatgaa atgntatggt tcnagtncct 420
 tttngggaca aattttctaa nctttttccn tttacnccnn tatnnggttn gttncaaatc 480
 ttttgggtcc ccttttctn naaancnccn atcatnttt ngntganttt tatncccca 540
 aantnaanna nncgctttt tttnttttt 570

<210> 2689
 <211> 566
 <212> nucleic acid
 <213> Glycine max

<400> 2689
 agaggttnnn ttttttnnat nnaaagtatt tttnnnnnnn nnnnnnnnnn nnnnnnnnn 60
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnngttagg naaatgacca 120
 accttagatc tnnaccacc taccngggtg ctttggtat naagggccna antggaggga 180
 cottgaaaag gctggcatca ctgttatcca aattgatgaa actgctttga gagaaagggt 240
 ctgccactga gggaaatcaa gaacaaggct cacttacttg gactgggctn gtncatgcct 300
 ttnagnaatc accaatgttg gnntntccn ngataccact caaaattcac accacatgt 360
 gctanctcca aacttttaac gacattnttc aatccatnaa ttnacatgg gcccttattg 420
 ttattcacca tttgaanaac tnttgcttcc gaattaanaa acttcctgt naaagtcntt 480
 tcntggaaaa ggtgttaaaa nattgggtct tgggaaattn ggncccttgn ngtttattta 540
 aaatnccant tccccanaa atnccc 566

<210> 2690
 <211> 300
 <212> nucleic acid
 <213> Glycine max

<400> 2690
 togcattcac gcgtacgtaa gctcggaatt cggctcgagc aaagaccctt gatttgataa 60
 agcaaggntt tccatctgga aaatttcttt ttgctggtgt tgttgatgga agattnattt 120

gggccaataa ttttgcattct tctctgaaca ccttcaggc acttgagac attgttgaa 180
atgacaaggt tgtggtttcc acgtcgtgtt ctcttcttca cactgcagtt gatctggtga 240
atgagaccaa attggaccaa gagattaagt cttggcttgc atttgcagca caaaaagttg 300

<210> 2691
<211> 239
<212> nucleic acid
<213> Glycine max
<400> 2691

ngtcgcangc acgcgtacgt nagctcgga ttcggctcga gcttgatttn atcaagcaag 60
gatttccatc tggaaaattt ctntncgctg gtgttggtga tngaaagana tatttggggc 120
aatagtcttg catcttctct gaacaccctt caggcacttg gggacattgt tgggaatgac 180
aaggttgtgg ttccacgtc ctgttctctt cttcacactg cagttgatct gtgaatgag 239

<210> 2692
<211> 303
<212> nucleic acid
<213> Glycine max
<400> 2692

tgcgatgcac gentaagtna gctcggaatt cggctcgagc atttactgcc aatgggtggg 60
tgcaatcata tggatccgc tgtgtcaaac ctcccatcat ctatggtgat gtgagccgtc 120
ccaagcccat gacagttttc tggctcttcaa ctgctcaaag ttgacccaaa cgaccaatga 180
agggaatgct tactggccct gttactattc tgaactggtc ctttgttaga gatgaccaac 240
caagatttga aacatgttac cagattgctt tggctatcaa ggatgagggt gaggatcttg 300
aga 303

<210> 2693
<211> 306
<212> nucleic acid
<213> Glycine max
<400> 2693

gtgcacgca cgcgtacgtn agctcggaat tccgctcgag ctgtgtcaag cctcccatca 60
tctatggtga tgtgagccgt cccaancca tgactgtttn ctggtcttnc aactgctcaa 120

agtttgacca aacgaccaat gaagggaatn cttactggcc ctgttactat tctgaactgg 180
 tcctttgtta gagatgacca gccaaagattc gaaacatgct accagattgc tttggctatc 240
 aaggatgagg ttgaggatct tgagaaagca ggtattactg tcatccagat tgatgaagct 300
 gctctt 306

<210> 2694
 <211> 459
 <212> nucleic acid
 <213> Glycine max
 <400> 2694

ccacgcgtcc ggtgttctct tcttcacact gcagttgacg tggatgaatga gaccaaattg 60
 gaccaagaga ttaagtcttg gcttgcatct gcagcacaaa aagttgttga agtaaatgcc 120
 ttggccaagg cattgtcttg acagaaggat gaggttttct tttctgctaa tgctgctgcc 180
 ttggcttcaa ggaagtcctc cccaagggtg ataaatgagg ctgtccaaaa agcgcgtgct 240
 gctctgaagg gctctgatca tcggagggcc acaaattgta gtgccagggt ggatgctcaa 300
 cagaagaaat tgaatctttc tgttcttcca acaactacaa ttggatcttt ccctcaaaact 360
 gccgatctta gaagagttcg tcgtgaattc aaggctaaca agatctccga ggaagattat 420
 atcccgtttc attaaggagg aaatttacao tgttgtaaa 459

<210> 2695
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2695

gcncgnacgc gtacgtaagc tcggaattcg gctcgagggt tttcttttct gctaattgctg 60
 ctgccttggc ttcaaggaag tcctcccaa gggtgataaa tgaggctgtc caaaaagccg 120
 ctgctgctct gaagggtctt gatcatcgga gggccacaaa tgtagtgcc aggttggatg 180
 ctcaacagaa gaaattgaat ctttctgttc ttccaacaac tacaattggn tctttccctc 240
 aaactgccga tcttagaaga gttcgtcgtg aattcaaggc taacaagatc tccgaggaag 300
 attata 306

<210> 2696

<211> 285
 <212> nucleic acid
 <213> Glycine max

 <400> 2696

 acgtcgcatg cacgcgtacg taagctcgga attcggctcg aggtaaatgc cttggccaag 60
 gcattgtctg gacagaagga tgaggttttc tttctgcta atgctgctgc cttggcttca 120
 aggaagtcct cccaagggt gataaatgag gctgtccaaa aagccgctgc tgctctgaag 180
 ggctctgata atcggagggc cacaaatgtt agtgccaggt tggatgctca acagaagaaa 240
 ttgaatcttt ctgttcttcc aacaactaca attnggtott tccct 285

<210> 2697
 <211> 303
 <212> nucleic acid
 <213> Glycine max

 <400> 2697

 acgtcgcatg cacgcgtacg taagctcgga attcggctcg agngaagctg tccaaaaggc 60
 cgtgctgct ctgaagggt ctgatcatcg gagggccaca aatgttagtg ccaggctgga 120
 ttctcaacag aagaaactga atcttctgt tcttccaaca actacaattg ngctctttcc 180
 ctcaaactgc cgatcttaga agagttcgcc gtgaattcaa ggctaacaag atctccgagg 240
 aanntatata catttcatta aggaggga ttaacaatgt tgtgaagctc caggaagaat 300
 tga 303

<210> 2698
 <211> 260
 <212> nucleic acid
 <213> Glycine max

 <400> 2698

 gtcgcntgca cgcgtacgta agctcggaat tcggctcgag tggatgaatga naccaaattg 60
 gaccaagaga ttaagtcttg gcttgcatth gcagcacaaa aagttgttga agtaaattgcc 120
 tggccaaggc attgtctgga agaaggatga ggttttcttt tctgctaattg ctgctgcctt 180
 ggcttcaagg aagtcctccc caagggtgat aaatgaggct gtccaaaaag ccgctgctgc 240
 tctgaagggc tctgatcatc 260

<210> 2699
 <211> 193
 <212> nucleic acid
 <213> Glycine max

 <400> 2699

 taacnagacg tcgcangcac gcgtacgtaa gctcgggaatt cggctcgagn tgataaatga 60
 ggctgtccaa aaagccgctg ctgctctgaa gggctctgat catcggaggg ccacaaatgt 120
 tagtgccagg ttggatgctc aacagaagaa attgaatctt totgtttctc caacaactac 180
 aattggatct ttc 193

<210> 2700
 <211> 307
 <212> nucleic acid
 <213> Glycine max

 <400> 2700

 gtcgcangca cgcgtacgta agctcgggaat tcggctcgag nctggtatta ctgtcatcca 60
 gattgatgag gctgctctaa gagaagggtt accctcgagg aagtctgagg aggttttcta 120
 tctaaactgg gctgttctact catttaggat taccaactgt ggtgtggagg aactactca 180
 gattcacact cacatgtgct actccaactt caatgacatc attcactcan tontagacat 240
 ggatgctgat gtgatcacca ttgagaactc cagatcagat gagaagttac tttcgggtctt 300
 ccgtgng 307

<210> 2701
 <211> 361
 <212> nucleic acid
 <213> Glycine max

 <400> 2701

 gtcgcatgca gcgtacgtaa gctcgggaatt cggctcgagg attcgaaaca tgctaccaga 60
 ttgctttggg ctatcaagga tgaggttgag gatcttgaga aagcaggtat tactgtcatc 120
 cagattgatg aagctgctct aagagaaggt ttacctctga ggaagtctga ggaggctttc 180
 tatctaaact gggctgttca ctcathtagg attaccaact gtggtgtgga ggacactact 240
 cagattcaca ctacatgtg ctactccaac ttcaatgaca tcatcactca atcatagaca 300

tggatgctga tgtgatcacc atgagaactc tagatcagac gagaagttac tttcagtctt 360

c 361

<210> 2702

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2702

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gctgctctaa gagaagggtt 60

acctctgagg aagtctgagg aggccttctt tctaaactgg gctgttcact catttaggat 120

taccaactgt ggtgtggagg acactactca gattcacact cacatgtgct actccaactt 180

caatgacatc attcactcaa tcatagacat ggatgctgat gtgatcacca ttgagaactc 240

tagatcagac gagaagttac tttcagtctt ccgcgaggga gtgaaatatg gtg 293

<210> 2703

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2703

gtcgcangca cgcgtacgta agctcggaat tcggctcgag attactgtca tccagattga 60

tgaagctgct ctaagagaag gtttacctct gaggaagtct gaggaggctt tctatctaaa 120

ctgggctggt cactcattta ggattaccaa ctgtggtgtg gaggacacta ctcagattca 180

cactcacatg tgctactcca acttcaatga catcattcac tcaatcatag acatggatgc 240

tgatgtgatc accattgaga actctagatc agacgagaag tt 282

<210> 2704

<211> 272

<212> nucleic acid

<213> Glycine max

<400> 2704

cgtcgcangc ncgcgtacgt nagctcgga ttcggctcga ggagatgacc aaccaagatt 60

tgaaacatgt taccagattg ctttggtat caaggatgag gttgnggatc ttgagaaagc 120

tggtattact gtcattcaga ttgatgaggc tgcncctaaga gaaggtttac ctctgaggaa 180

gtctgaggag gctttctatc taaactgggc tgttactca tttaggatta ccaactgtgg 240
 tgtggaggac actactcaga tncacactca ca 272

<210> 2705
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2705

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag aagaagttca accttccaat 60
 cctcccaacc accacaattg gatccttccc tcagactggt gaactgagga ggggtcgctcg 120
 tgaatacaag gctaacaaga tctcagagga ggagtatggt agttcaatta aagaggaaat 180
 ccgcaaagtt gttgaactcc aagagaatct tgatatcgat gtcctggtag acggggagcc 240
 tgagaggaat gacatggtgg agtactttgg tgagcagttg tcaggctttg cctttacc 298

<210> 2706
 <211> 307
 <212> nucleic acid
 <213> Glycine max
 <400> 2706

nnogcatgca cgcgtacgta agctcggaat tcggctcgag aagaagttca accttccaat 60
 cctcccaacc accacaattg gatccttccc tcagactggt gaactgagga ggggtcgctcg 120
 tgaatacaag gctaacaaga tctcagagga ggagtatggt agttcaatta aagaggaaat 180
 ccgcaaagtt gttgaactcc aagagaatct gcatatcgat gtcctggtag acggngagcc 240
 tgagaggaat gacatggtgg agtactttgg tgagcagttg tcaggctttg cctttaccgt 300
 taaggct 307

<210> 2707
 <211> 452
 <212> nucleic acid
 <213> Glycine max
 <400> 2707

ccacgcgtcc gctggaagca gatggctgat gctggaataa agtatattcc tagcaacacc 60
 ttctcacttt acgatcaagt actggacaca acagccatgc tcggggcagt tccatctaga 120

tataattgga atggtgggga gattggggtt gatgtttact tctcaatggc aagaggggaat 180
gcattctgtac cagctatgga aatgaccaag tggtttgaca ccaattacca ttacattggt 240
cctgaattgg gtctctgatgt taagttctcc tatgcatcac acaaggctgt cgatgaattt 300
aaagaggcca aagttctggg agttaatact gtacctgtgc ttgtgggacc tgtatcctac 360
ttgttgctgt caaaaccagc taagggtgtt gaagaagtca tttcccttc tttccctaatt 420
tgacaagatc cttcctgtct acagggangt tt 452

<210> 2708
<211> 260
<212> nucleic acid
<213> Glycine max

<400> 2708

acgtcgang cagcgtagc taagctcgga attcggtcgc agcaagtggg ttgacaccaa 60
ttncattac attgttctg aattgggtcc tgatgttaag ttctcctatg catcacacaa 120
ggctgtcgat gaatttaaag aggccaaagt tctgggagtt aatactgtac ctgtgcttgt 180
gggacctgta tctacttgt tgctgtcaaa accagctaag ggtgttgaga agtcattttc 240
ccttctttcc ctaattgaca 260

<210> 2709
<211> 275
<212> nucleic acid
<213> Glycine max

<400> 2709

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gcaagaggga attcatctgt accagctatg gaaatgacca agtggtttga cacaaactat 180
cattacattg ttctgaatt ggggccagat gttaagttct cctatnccat cacacaagggt 240
tgtggatgaa tacntngagg ctaaagttct gggaa 275

<210> 2710
<211> 304
<212> nucleic acid
<213> Glycine max

<400> 2710

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ttcgctctcg agtctttctg ggatggcaag agcagcgccg aggatttgca gaaggtgtct 180
tgtgatctca gggcatccat ctggaagcag atggctgatg ctgggatcaa gtacatcccc 240
agcaacactt tctctcacta tgaccagggt ctcgacgcca ccgccaccct cggtgccggt 300
ccac 304

<210> 2711

<211> 341

<212> nucleic acid

<213> Glycine max

<400> 2711

tcgcangcac gcgtacgtaa gctcggaatt cggtctgagc tccactcaga agcgaagaan 60
aagccacaga gaactagtct cctacttctc acccgcaaga naaaaaatggc atctcacatc 120
gttggatacc cccgatggg tcccaagaga gagctcaagt tcgctctcga gtctttctgg 180
gatggcaaga ncagcgccga ggatttgagc aaggtgtctt ctgatctcag ggcatccatc 240
tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt ctctcactat 300
gaccagggtc tcgacgccac cgccaccctc ggtgccgttc c 341

<210> 2712

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2712

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gaaccagtct cctactctct ctcaccacaca agaaaaatgg catctcacat cgttggatac 120
cccgcatgg gtcccaagag agagctcaag ttcgctctcg agtctttctg ggatggcaag 180
agcagcgccg aggatttgca gaaggtggct gctgatctca ggcatccat ctggaagcag 240
atggctggtg ctgggatcaa gtacatcccc agcaacactt tctcgttcta tgaccagctg 300
ctcgacgcca ccgccaccct cg 322

<210> 2713
 <211> 328
 <212> nucleic acid
 <213> Glycine max

 <400> 2713

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 ttctgatctc agggcatcca tctggaagca gatggctgat gctgggatca agtacatccc 240
 cagcaacaact ttctctcact atgacanggt tctcgacgcc accggcaccc tcggtgccgt 300
 tncaccaagt aggtggancc gcggcgag 328

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<210> 2714
 <211> 339
 <212> nucleic acid
 <213> Glycine max

 <400> 2714

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 catctcacat cgttggatac ccccgcatgg gtcccaagag agagctcaag ttcgctctcg 180
 agtctttctg ggatggcaag agcagcgccg aggatttgca gaagggtggct gctgatctca 240
 ggtcatccat ctggaagcag atggctgggt ctgggatcaa gtacatcccc agcaaacatt 300
 tctcgttcta tgaccagctg ctcgacgcca ccgccaccc 339

<210> 2715
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 2715

 catgcacgcg tacgtaagct cggaattcgg ctccgagcac agagaaccag tctcctactc 60
 tctctcacc cacaagaaaaa tggcatctca catcgttgga taccctcgca tgggtcccaa 120
 gagagagctc aagttcgctc tcgagctctt ctgggatggc aagagcagcg ccgaggattt 180

gcagaaggtg gctgctgac tcaggtcacc catctggaag cagatggctg gtgctgggat 240
 caagtacacc cccagcaaca ctttctcggt ctatgaccag ctgctcgacg ccaccg 296

<210> 2716
 <211> 308
 <212> nucleic acid
 <213> Glycine max
 <400> 2716

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 gatctcaggg catccatctg gaagcagatg gctgatgctg ggatcaagta catccccagc 120
 aacactttct ctactatga ccaggttctc gacgccaccg ccaccctcgg tgcggtccac 180
 caaggtaagg ctggcaccgg cggcgagatt gggtttgata cctacttcnc catggccaga 240
 ggtaatgcta ccgtgccagc tatggagatg accaagtggg tcgacaccaa ctaccacnct 300
 nttgtccc 308

<210> 2717
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 2717

anntnnnctg acgtaanctc ggaattcggc tcgagcttgc tccctcanan ncntgaanaa 60
 gccacagaga actagtctcc taactctcac ccgcaagacn aaaaatggca tctcacatcg 120
 ttggnatccc ccgtatnggt cccaagagag agcncaagtt cgctctcgag tctttctggg 180
 atggcaagag cagcgccgag gatttgaga aggtgtcttc tgatctcagg gcatccatct 240
 ggaagcagat ggctgatgct gggatcaagt acatccccag caacactttc tctactatg 300
 accaggttct cgac 314

<210> 2718
 <211> 307
 <212> nucleic acid
 <213> Glycine max
 <400> 2718

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctccctcaga agcgaagaag 60

aagccacaga gaaccagtct cctactctct ctcacccaca agaaaaatgg catctcacat 120
 cgttggatac ccccgcatgg gtcccaagan agagctcaag ttcgctctcg agtctttctg 180
 ggatggcaag agcagcgccg aggatttgca gaaggtggct gctgatctca ggcatccat 240
 ctggaagcag atggctgggtg ctgggatcaa gtacatcccc agcaacactt tctcgttcta 300
 tgaccag 307

<210> 2719
 <211> 318
 <212> nucleic acid
 <213> Glycine max
 <400> 2719

acgcgtacgt aagctcggaa ttcggctcga ggcttcaact tgctccctca gaagcgaaga 60
 agaagccaca gagaaccagt ctctactct cttcaccca caagaaaaat ggcatctcac 120
 atcgttggat acccccgcat ggggtccaag agagagctca agttcgctct cgagtctttc 180
 tgggatggca agagcagcgc cgaggatttg cagaaggtgg ctgctgatct caggatccatcc 240
 atctggaagc agatggctgg tgctgggatc aagtacatcc ccagcaacac tttctcgttc 300
 tatgaccact gctcgacg 318

<210> 2720
 <211> 324
 <212> nucleic acid
 <213> Glycine max
 <400> 2720

gcacgcgtac gtaagctcgg aattcggtc gagctcagaa nogaanaaga agccacncna 60
 gaactagtct cctactctca cccgcaanaa aaaaatggca tctcacatcg ttngatacc 120
 cccgtatggg tccaagaga gagctcaagt tcgctctcga gtctttcttg gatggcaaga 180
 gcagcgccga ggatttgca aaggtgtctt ctgatctcag ggcatccatc tggaagcaga 240
 tggctgatgc tgggatcaag tacatcccca gcaacacttt ctctcactat gaccagggtc 300
 tcgacgccac cgcaccctc ggtn 324

<210> 2721
 <211> 331
 <212> nucleic acid

<213> Glycine max

<400> 2721

nnncnngca tncgtacgta agctcggaat tcggctcgag cggctcgagt tcaacttget 60
ccctcagaag cgaagaagaa gccacagaga actagtctcc tactctcacc cgcaagaaaa 120
aaatggcatc tcacatcggtt ggataccccc gtatgggtcc caagagagag ctcaagtctg 180
ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagnag gtgtcttctg 240
atctcagggc atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 300
acactttctc tcactatgac caggttctcg n 331

<210> 2722

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2722

gttgcangca cgcgtacgta agctcngaatt tcggctcgag gccacagaga ctccagtctc 60
ctactctctc tcaccanana gananntggn atntcacntc gttggatacc cccgcatggg 120
tcccaagana gagctcnagt tcgctctcga gtctttctgg gatggcnagn nongcgccga 180
ggatttggca naaggtggct gctgatctca ggtcatccat ctogaagcag atggctggtg 240
ctgggatcaa gtacntcccc agcaacactt tctcgttcta tgaccagctg ctogacgccca 300
ccgccaccct cggtgccgtc cccccag 327

<210> 2723

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 2723

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acatccccag caacactttc tctcactang accnggtttc tcgacgccac cgcnacccctc 120
ggtgccgttc caccaaggna cggctggacc ggcggcgaga ttgggtttga tacctacttc 180
tccatggcca nangtaatgc taccgtgcca gctatggaga tgaccaagtg gttcganacc 240
aactancact ttattgtccc tgaatgggcn ctgatgtgaa cttcacctat gcttctcaca 300

aggcngctga tgaana

316

<210> 2724

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2724

cgntnnnaag cgtacgttag ctcggaattc ggctcgagct tgctccctca gaagcgaaga 60

agaagccaca gagaaccagt ctctactct ctctaccca caagaaaaat ggcatctcac 120

atcggttgat acccccgcat ggggcccaag agagagctca agttcgctct cgagtcttct 180

gggatggcaa gaggcgcgc gaggatttgc agaaggtggc tgctgatctc aggtcatcca 240

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atgaccagct gctcgacgcc 320

<210> 2725

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2725

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctgagaagcg aagaagaagc 60

cacagagaac tagtctcta ctctacccg caagaaaaaa atggcatctc acatcggttg 120

atacccccgt atgggtccca agagagagct caagttcgct ctcgagtctt tctgggatgg 180

caagagcagc gccgaggatt tgcagaaggt gtcttctgat ctgagggcat ccatctggaa 240

gcagatggct gatgctggga tcaagtacat cccagcaac actttctctc actatgacca 300

g 301

<210> 2726

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2726

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gagaaaaatg gcatctcaca tcgttagata ccccgcatg ggncccaaga gagagctcaa 120

gttcgctctc gagtctttct gggatggcaa gaggcagccc gaggatttgc agaaggtggc 180
 tgotgatctc aggtcatcca tctggaagca gatggctggt gctgggatca agtacatccc 240
 cagcaaacct ttctcgttct atgaccagct gctncgacgc caccgccacc ctgggtgccg 300
 tccccccan gt 312

<210> 2727
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 2727

nnogtctcan gcacgcgtac gtnagctcnn naattcggct cgaggaacta gtctcctact 60
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 tgcagaaggt gtcttctgat ctccagggcat ccacatctggaa gcagatngct gatgctggga 240
 tcaagtacat cccanacaac actttctctc actatgacca ggtncctgac gccnccgcc 300
 t 301

<210> 2728
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2728

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 atggcatctc acatcgttgg atacccccgc atgggtccca agagagagct caagttcgct 180
 ctcgagtctt tctgggatgg caagagcagc gccgaggatt tgcagaaggt ggctgctgat 240
 ctccaggtcat ccacatctggaa gcagatggct ggtgctggga tcaagtacat cccagcaac 300
 actttctcgt tctatg 316

<210> 2729
 <211> 330
 <212> nucleic acid
 <213> Glycine max

<400> 2729

natctcangc aagcgtacgt attctcgnna ttnggcnacg aggccttcaac ttgctccctc 60

agaancgaag aagaagnnac agananctan tctcctactc tcacccgcaa gaanaaaatn 120

gcattctcaca tcgttgata ccccgatg ggtcccaaga gagagctcaa gttcgtctc 180

gagtctttct gggatggcaa gagcagcgn gaggatttgc agaaggtgtc ttctgatctc 240

aggcatcca tctggaagca gatggctgat gctgggatca agtacatccc cagcaacact 300

ttctctcata tgcaccaggt tctcgaagcc 330

<210> 2730

<211> 271

<212> nucleic acid

<213> Glycine max

<400> 2730

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gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc tggaagcaga 120

ggctgatgct gggatcaagt acatccccag caacactttc tctcactatg accaggttct 180

ngacgcnacc gncannctcg gtgntgtnc c atnaaggnan gnnatatann gttgtnatnn 240

nggntnnann ntnttttcca ngttntgcag g 271

<210> 2731

<211> 318

<212> nucleic acid

<213> Glycine max

<400> 2731

tcccgtagct nagctcgga ttcggctcgn ggcttcanct tgctccctca gaagcgaaga 60

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cgttgatgac ccccgatgc gtcccaagan agagctcaag ttcgctctcg agtctttctg 180

ggatgcaaga gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc 240

tggaagcaga tggctgatgc tgggatcaag tacatcccc gcaacacttt ctctcactat 300

gaccaggttc tcgacgca 318

<210> 2732

<211> 307
 <212> nucleic acid
 <213> Glycine max

 <400> 2732

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 aaatggcatc tcacatcggt ggataccccc gtatgggtcc caaganagag ctcaacttcg 180
 ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg 240
 atctcagggc atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 300
 acacttt 307

<210> 2733
 <211> 304
 <212> nucleic acid
 <213> Glycine max

 <400> 2733

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 atggcatctc acatcggttg atacccccgc atgggtccca agagagagct caagtctgct 180
 ctcgagtctt tctgggatgg caagagcagc gccgaggatt tgcagaaggt ggctgctgat 240
 ctcaggtcat ccatctggaa gcagatggct ggtgctggga tcaagtacat ccccagcaac 300
 actt 304

<210> 2734
 <211> 333
 <212> nucleic acid
 <213> Glycine max

 <400> 2734

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 acccacaaga naaatgncat ctcacatgcn tggatanccc cgcattgggtc ccnaganaga 180
 gctcaagttc gcnctcgagt ctttctggga tggcaagagc agcgccgagg atttgcagaa 240

ggtggctgct gatctcaggt catccatctg gaagcagatg gctgggtgctg ggatcaagta 300
catccccagc aacactttct cgttctatga cca 333

<210> 2735
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 2735

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agaagaagcc acagagaact agtctcctac tctcaccgc aagaaaaaaaa tggcatctca 120
catcgttgga tcccccgta tgggtcccaa gagagagctc aagttcgctc tcgagtcttt 180
ctgggatggc aagagcagcg ccgaggattt gcagaagggtg tcttctgata tcagggcatc 240
catctggaag cagatggctg atgctgggat caagtacatc ccagcaaca ctttctctc 299

<210> 2736
<211> 333
<212> nucleic acid
<213> Glycine max

<400> 2736

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gccacagaga actagtctgc ctactctcac ccgcaagana aaaatggcat ctacatgcg 120
ttggataccc ccgtatgggt cccaagagag agctcaagtt cgctctcgag tctttctggg 180
atggcaagag cagcgccgag gatttgaga aggtgtcttc tgatctcagg gcatccatct 240
ggaagcagat ggctgatgct gggatcaagt acatccccag caacactttc tctactatg 300
accaggttct tcgacgcnac gccacctcgg tgc 333

<210> 2737
<211> 320
<212> nucleic acid
<213> Glycine max

<400> 2737

agtcgcangc acgcgtacgt aagctcgnaa attcggctcg agttgctccc tcagcagcna 60
agaagaagcc acagagaact agtactcnct antctcacc gcaagannna aatggcatct 120

cacatcgntg gatacccccg tatgggtccc aagagagagc tcnagttcgt nctcgagtct 180
 ttctgggatg gcaagagcag cgccgaggat ttgcagaagg tgtcttctga tctcagggca 240
 tccatctgga agcagatggc tgntgctggg atcaagtaca tccccagcaa cactttctct 300
 nactatggcc agttctcnac 320

<210> 2738
 <211> 287
 <212> nucleic acid
 <213> Glycine max
 <400> 2738

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 gaagccacag agaaccagtc tcctacttct ctctcaccca caagaaaaat ggcatctcac 120
 atcgttggat acccccgcat ggggtccaag agagagctca agttcgtctt cgagtctttc 180
 tgggatggca agagcagcgc cgaggatttg cagaaggtgg ctgctgatct caggtcaccc 240
 atctggaagc agatggctgg tgctgggatc aagtacatcc ccagcaa 287

<210> 2739
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2739

gttgcatgtn agcgtacgta agcttcggaa ttctggtnnn ggatttggat cgagcttnnt 60
 ccctcagnag cgnagaagaa gccncagaga actagtctcn tactctcacc cgcaagaaaa 120
 aaatggcatc tcacatcgtt ggatacccc gtatgggtcc caagagagag ctcaagttcg 180
 ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg 240
 atctcagggc atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 300
 acactt 306

<210> 2740
 <211> 291
 <212> nucleic acid
 <213> Glycine max
 <400> 2740

gtcgcangca cgcgtagcta agctcggaat tcggctcgag aacttgctcc ctcagaagcg 60
aagaagaagc cacagagAAC cagtctccta ctctctctca cccacaagaa aaatggcatc 120
tcacatcggt ggatacccc gcattgggtcc caagagagag ctcaagttcg ctctcgagtc 180
tttctgggat ggcaagagca gcgccgagga tttgcagaag gtggctgctg atctcaggtc 240
atccatctgg aagcagatgg ctggtgctgg gatcaagtac atccccagca a 291

<210> 2741
<211> 322
<212> nucleic acid
<213> Glycine max
<400> 2741

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gcacgcgtac gtaagctcgg aattcggctc gagatcttgc tccctcagaa gogaagaaga 60
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ggatggcaag cgcagcgccg aggatattgca gaaggtgtct tctgatctca gggcatccat 240
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tgaccagttc tcgacgccac gn 322

<210> 2742
<211> 310
<212> nucleic acid
<213> Glycine max
<400> 2742

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agccacagag aactagtctc ctactctaca cccgcaagaa aaaaatggca tctacacatc 120
gttggatacc cccgataggg tcccaagaga gagctcaagt tcgctctcga gtctttctgg 180
gatggcaaga gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc 240
tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaaacattt ctctcatang 300
accaggttct 310

<210> 2743
<211> 304
<212> nucleic acid

<213> Glycine max

<400> 2743

agttgcangc acgcgtacnt aagctcggaa ttcggctcga gtcagaagcg aagaagaagc 60
cacagagaac tagtcnnnt actctcacc cgaagaaana aatngccatc tcanatgcgt 120
tggatncccc cgtatgggtc ccaagagaga gctcaagttc gctctcgagt ctttctggga 180
tggcaangnc ancgccgagg atttgcagaa ggtgttttct gatctcaggg catccatctg 240
gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct ntcactatga 300
ccag 304

<210> 2744

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2744

tcgcangnac nntcggaaat tcggctcgag cttgctccct cagaagcgaa gaagaagcca 60
cagagaacca gtctcctact ctctctcacc cacaagaaaa atggcatctc acatcggttg 120
atacccccgc atgggtccca agagagagct caagttcgct ctcgagtctt tctgggatgg 180
caagagcagc gccgaggatt tgcagaaggt ggctgctgat ctcaggatcat ccactctggaa 240
gcagatggct ggtgctggga tcaagtacat cccagc 277

<210> 2745

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2745

gtngcangca cgcgtacgta agctcggaaat tcggctcgag ctccctcaga agcgaagaag 60
aagccacaga gaactagtct cctactctca cccgcaagan aaaaatggca tctcacatcg 120
ttggataccc ccgtatgggt cccaagagng agctcaagtt cgctctcgag tctttctggg 180
atggcaagag cagcgccgag gatttgcaga aggtgtcttc tgatctcagg gcatccatct 240
ggaagcagat ggctgatgct gggatcaagt acatccccag caacactt 288

<210> 2746

<211> 318
 <212> nucleic acid
 <213> Glycine max

 <400> 2746

 gttgcangca cgcgtacgta agctcgggaat tcggctcgag cagaagcgaa gaagaagcca 60
 nagagaacta gtctcctacn nttcaccgcg aagaaaaaat ggnatctcac atcgttggat 120
 acncccgat gggtnnccaa gagagngnna agttcgntct cgagtccttc tgggatggca 180
 agagcagcgc cgaggatttg cagaaggtgt cttctgatct cagggcatcc atctggacgc 240
 agatggctga tgctgggatc aagtacatnn ncagcaanac tttctctcan tatgaccagg 300
 ttctcgacgc naccgcca 318

<210> 2747
 <211> 331
 <212> nucleic acid
 <213> Glycine max

 <400> 2747

 gcaaantcgc tgnangcnta cgtncgctcg ganttcggat ngagctcgag ccgctgcttc 60
 aacttgctac ctcagaagcg aagaagaagc cacagagaac cagtctccta ctctctctca 120
 cccacaagaa naatgggatc tcacatcggt ggataccccc gcatgggtcc caagagagag 180
 ntcnagttcg ctctcgagtc tttctgggat ggcaagngcn gcgccgagga tttgcagaag 240
 gtggctgctg atctcaggtc atccatctgg aagccagatg gctgggtgctg ggatcaagta 300
 catccccagc aacactttct cgtttatgnc c 331

<210> 2748
 <211> 307
 <212> nucleic acid
 <213> Glycine max

 <400> 2748

 aatcgcangc aagcgtacgt aagctcgga ttcggctoga gctctgcttc aacttgctcc 60
 ctcagaagcg aagaagaagc cacagagaac cagtctccta ctctctctca cccacaagaa 120
 aaatggcatc tcacatcggt ggataccccc gcatgggtcc caagagaagc tcaagttcgc 180
 tctcgagtct ttctgggatg gcaagagcag cgccgaggat ttgcagaagg tggctgctga 240

tctcaggtca tccatctgga agcagatggc tgggtgctggg atcaagtaca tccccagcaa 300
cattttct 307

<210> 2749
<211> 302
<212> nucleic acid
<213> Glycine max
<400> 2749

gtcgcangna cgcgtacgta agctcggaat tcggctcgag cttgntccct cagaagcgaa 60
gaanaancca cagagaanta gtctnctact ctgcacccgc aanaaaaaaa tggcntctca 120
catgcgttgg atacccccgt atgggtccca aganagagct caagttcgct ctcgagtctt 180
tctgggatgg caagagcagc gcgaggatt tgcagaaggt gtcttctgat ctcagggcat 240
ccatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac actttctctc 300
an 302

<210> 2750
<211> 287
<212> nucleic acid
<213> Glycine max
<400> 2750

cangcacgcg tacgtnagct cggaattcgg ctcgagnctt gctccctcag aagcgaagaa 60
gaagccacag agaaccagtc tctactctc tctcaccac aaganaaatg gcattctcaca 120
tcgttgata cccccgatg ggtcccaaga nagagctcaa gttegtctctc gagtncttct 180
gggatggcaa gagcagcgcc gaggatttgc agaaggtggc tntgatctc aggtcatcca 240
tctggnagca gatggctggg gctgggatca agtacatccc cagcaac 287

<210> 2751
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2751

agtencangc acgcgtacgt aagctcgga ttcggctcga gctggcttca anttgctccc 60
tcagaagcga agaagaagcc acagaganct ngctctctac tctcaccgc aaganaaaaa 120

tggcatctca catcgttggga tccccccgta tgggtcccaa ganagagctc aagttcgctc 180
 tcgagtcttt ctgggatggc aagagcagcg ccgaggnttt gcagaagggtg tcttctgntc 240
 tcagggcata catctggaag cagatggctg atgctgggat caagtacatc cccagcaaca 300

<210> 2752
 <211> 285
 <212> nucleic acid
 <213> Glycine max

<400> 2752

ncttncaagc acgcgtacgt nagctcggaa ttcggctcgg gctccctcag aagcgaagaa 60
 gaagccacag agaaccagtc tcctantcnc tctcaccac aagaaaaatg gcattctaca 120
 tcgttggata cccccgcatg ggtcccaaga nagagctcaa gtctgctctc gagtctttct 180
 gggatggcaa gagnagcgcc gaggatttgc agaagggtggc tgctgatctc aggtcatcca 240
 tctggaagca gatggctggg gctgggatca agtacatcnc cagcn 285

<210> 2753
 <211> 326
 <212> nucleic acid
 <213> Glycine max

<400> 2753

tgctgtaana nanaangtng cgtgcacgcn tacgtaaagc tcgggaattc ggctcgagct 60
 ggnttcaant tggntcctca naagcgaaga agaagccaca gagaaccagt ctctantnt 120
 ctntcacca caagaaaaat ggcntctcac atcgttggat acccccgcat gggcccaag 180
 anagagtca agttcgntnt cgagtctttc tgggatggca agancagcgc cgaggatttg 240
 cagaagggtg ctgntgatnt caggtcatcc atntggaagc agatggctgg tgntgggatc 300
 aagtacatcc ccagcaanac tttntc 326

<210> 2754
 <211> 281
 <212> nucleic acid
 <213> Glycine max

<400> 2754

ncgtcgcang cagcgtacg taagctcgga attcggctcg aggtccctc agaagcgaag 60

aagaagccac agagaaccag tctctactc tctctcacc acaagaaaaa tggcatctca 120
catcgttgga taccocccgca tgggtcccaa gagagagctc aagttcgctc tcgagtcctt 180
ctgggatggc aagagcagcg ccgaggattt gcagaaggtg gctgctgac tcaggtcatc 240
catctggaag cagatggctg gtgctgggat caagtacatc c 281

<210> 2755
<211> 303
<212> nucleic acid
<213> Glycine max
<400> 2755

gtcgcangca cgcgtncgac gantacgtna nctcggnntt nggntccncg ctnacatggg 60
catngnaaac tgtttacgag tgagtntcct angencancc gcaagaaaaa aatggcatct 120
cacatcggtg gatacccccg natgggttcc caagagagag ctcaagttcg ctctcgagtc 180
tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg atctcagggc 240
atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca acatttctct 300
cat 303

<210> 2756
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2756

cgtcgcatgc acgcgtacgt nagctcggnn ttcggctcgn gcggctcng gcttcaactt 60
gctccctcag aagcgaagaa gaagccacag agaactagtc tctaacctc acccgcaaga 120
naaaaatggc atctcacatc gttggatacc cccgtatggg tcccaagaga gagctcaagt 180
tcgtctcga gtctttcttg gatggcaaga gcagcgccga ggatttgcag aaggtgtctt 240
ctgatctcag ggcattccatc tggaagcaga tggctgatgc tgggatcaag tacatcccca 300

<210> 2757
<211> 299
<212> nucleic acid
<213> Glycine max
<400> 2757

tngcangcac gcgtacgtaa gctcggaatt cggctcgagc tgcttcaact tgctccctca 60
gaagcgaaga agaagcnaca gaggnccagt ctctactct ctctaccca caagaaaaat 120
ggcatctcac atcgttggat acccccgcat gggteccaag agagagctca agttcgctct 180
cgagtctttc tgggatggcn ngagcagcgc cgaggatntg cagaaggtgg ctntgatct 240
caggncatcc atctggaagc aganggctgg tgcngggatc aagtacatcc ncagcnacn 299

<210> 2758
<211> 310
<212> nucleic acid
<213> Glycine max

<400> 2758

tnncatgcac gcgtacgtaa gctcggaatt cggctcgagg ctccctcnga agcgaagaag 60
aagccacaga gaactagtcn nctantctca cccgcaagaa naaaatgngc atcttcacat 120
ncgttggata ccccgctatg ggtcccaaga gaganctcaa gttecgctctc gagtctttct 180
gggatggcaa gaggcagcnc gaggatttgc agaaggtgtc ttctgatctc agggcatcca 240
tctggaagca gatggctgat gctgggatca agtacatccc cagcaacact ttctnccan 300
tntgaccagg 310

<210> 2759
<211> 318
<212> nucleic acid
<213> Glycine max

<400> 2759

acgcatgnag cgtacgnag ctcggnattc ggctcgagct gcttcaactt nctccctcag 60
aagcgaagaa gaagccacag agaactantc tctactctc acccgacta aaanaatggc 120
atctcacatc gttggatacc cccgtatngg ncccaagana gagctcaagt tcgctctcga 180
gtctttcttg gatngcnnga ncagngccgn ggatttgcag caggtgtctt ctgatctcan 240
ggcatccatc tggnnacaga tggctgntgc tgggatcaag tncatccnca ncaacacttt 300
ctctcactag nncaggtt 318

<210> 2760
<211> 293
<212> nucleic acid

<213> Glycine max

<400> 2760

nncgtcgcan gcacgcgtac gtaagctcgg taattcggct cgagnaactt gctccctcag 60
aagcgaagna gaagccacag agnactagtc tcctacttct caccgcgnag anaaanntgg 120
catctcacat cgttggatac ccccgatatg gtcccaagag agagctcaag tncnctctcn 180
agtctttctg ggatggcnag agcagcgccg aggatttgca gaaggtgtct nctnatctca 240
gggcatccat ctggaagcng atggctgatg ctgggatcaa gtacatcccc agc 293

<210> 2761

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2761

ngntcangca cncgtacgtn agctcggaat tcnctcgan gctccctcag aagcgaagaa 60
nnagccacag agaactngnc tcctactctc acccgcaaga aaaaaatggc acctcanntc 120
gttggatacc cccgatgng tcccaagagg gagctcaagt tcgctctcga gtcttctggg 180
atggcaagag cagcgccgag gatttgca gaaggtgtctc tgatctcagg ggcattccatc 240
tggaagcaga tggctgatgc tgggatcaag tacatcccc gcancacttt ctctcactat 300
gn 302

<210> 2762

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2762

gatgctngca cgcgtacgtn agctcggaat tnggctcgag cagagaacta gtctcctact 60
ctcaccgcga agaaaaaat ngatctcac atcgttggat acccccgat gggcccaag 120
agagagcnca agttcgctct cgagttcttt ctnggatggc aagngcagct ccgaggattt 180
gcagaaggtg tcnnntgatc tcagggcctc catctggaag cagatggctg atgctgggat 240
caagtacatc ccngcaaca ntttctcna ctctgacaa ggttctcgac gcnaccgcga 300
accctcgggtg n 311

<210> 2763
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2763

 cncgtacgta agctcggaat tcggctcgag cnaagaanaa gccacagcag anccagtctc 60
 ctantctctc tcanccacaa nanaaatngc atencacatc gtnggatacc cccgcatggn 120
 taccceanag agagnncaag ttacgctctc gagtctttct gggatggcaa gagcagcgcc 180
 gaggatttgc agaaggtggc tgctgntctc aggtcatcctn tctggaagca gatggctggt 240
 gctgggatca agtacatccc cancaacact ttctcgttct atgaccagct gctcgacg 298

<210> 2764
 <211> 300
 <212> nucleic acid
 <213> Glycine max

 <400> 2764

 tcnangcacg cgtacgtnaa gctcggaatt tcggaattcg gctcgagact tgctccctca 60
 gaagcgaaga agaagccaca gagaaccagt ctctactctc ctctaccca caagaaaaat 120
 ggnatctcac atcgttggat acccccgcac gggncccaag agagagctcn agttcgntct 180
 cgagtctttc tgggatggna anagcancgn cgangntttg canaangngg ctgctggtct 240
 cangncatcc atctggaanc ngatggctgg tgctgggatc nagtacatcc ccagcaacac 300

<210> 2765
 <211> 330
 <212> nucleic acid
 <213> Glycine max

 <400> 2765

 gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gcttcaactt gctccctcag 60
 aagcgaagaa gaagccacag agaactagtc tccttacttc tcacccgcaa ganaaaaaatg 120
 ggcattctcac atcgttggat acccccgtat gggcccaag anagagctca agttcgctct 180
 cgagtctttc tgggatggca agagcagcgc cgaggatttg cagaagggtg cttctgatct 240
 caggcatcc atctggaagc agatggctga tgctgggatc aatacatccc cagcaacatt 300

tctctcatat gaccagttct cggacgccac

330

<210> 2766
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 2766

ttgangcacg cgtacgtnag ctcggaattc ggetcgagct tcaacttgct ccctcagang 60

cgaagaagan gccacagaga acnagtctcc tactctcacc cgcaaganaa aaatggcatc 120

tccacatcgt tggatacccc cgtatgggtn cccaaganag agtcaagtt cgctctcgag 180

tctttctggg atggcaagag cagcgccgag gatttgcaga aggtgtcttc tgatctcagg 240

gcatccatct ggaagcagat ggctgatgct gngantcaag tacatcccca gcaacacttt 300

ctctcact 308

<210> 2767
<211> 309
<212> nucleic acid
<213> Glycine max

<400> 2767

gttnaatgca cgcgtacgta agtcggaat tcggctcgag gctccctcag aagcgaagaa 60

gaagccacag agaaccagtc tcctactctc tctcaccac aanaaaaatg gcatctcaca 120

tcgttgata ccccgcatg ggtcccaaga gagagctcaa gtctgctctc gagtctttct 180

gggatggcaa gagcagcgc gaggatttgc agaagggtggc tgctgatctc aggttcatcc 240

atctggaagc agatggctgg tgctggggat caagtacatc cccagcaaca cttctcgttc 300

tatgaccag 309

<210> 2768
<211> 247
<212> nucleic acid
<213> Glycine max

<400> 2768

ctanaactnn acnccctcag aagcgaagaa gaagccacag agaactagtc tcctactctc 60

acccgcaaga naaaaatggc atctcacatc gttggatacc cccgtatggg tcccaagaga 120

gagctcaagt tcgtctcgag tcttctggg atggcaagag cagcgccgag gatttgcaga 180
 aggtgtcttc tgatctcagg gcatccatct ggaagcagat ggctgangct gggntcaagt 240
 acatccc 247

<210> 2769
 <211> 248
 <212> nucleic acid
 <213> Glycine max
 <400> 2769

gcacgcgtac gtaagctcgg aattcggctc gagctccctc agaagcgaag aanaagccac 60
 agagaactag tctnctactc tcannccgca agaaaaaaat ggcattctcac atcgttggat 120
 acccccgtat ggggtcccaag agagagctca agtctcgtct cgagtctttc tgggatggca 180
 agagcagcgc cgaggatttg cagaaggtgt cttctgatct cagggcatcc atctggaagc 240
 agatggct 248

<210> 2770
 <211> 284
 <212> nucleic acid
 <213> Glycine max
 <400> 2770

ncgttgcang cacgcgtacg tnagctcgga attnngctcg agtctgcttc aacttgctcc 60
 ctcagaancg aagaagaagc cacagaganc tagtctccta ctctcaccgc caagaaaaaa 120
 atggcatctc acatcgttgg ntacccccgt atgggtccca agaganagct caagtctgct 180
 ctcgagtctt tctgggatgg cnagngcagc gccgaggatt tgcagaaggt gtcttctgat 240
 ctcagggcat ccattctggaa gcagatggct gangctggga tcan 284

<210> 2771
 <211> 329
 <212> nucleic acid
 <213> Glycine max
 <400> 2771

anaancaatg caccgtacnt aantcggntc natgncgagc ngaattcggc tcgagctctg 60
 cttcaacttg ctcccatnga agcgaanaag aagccacaga nnactantct cctantctca 120

cncgcaagan aanaatggca nctcacatgc gttggataacc ccntatggg tcccaagaga 180
gagctcaagt tcgctctcga gtctttctgg gatngcaaga gnagcgccga ggatttgcn 240
aaggngtctt ctgatctcag ggcattccatc tggnagcana tggctgatgc tnggancaag 300
tacatcccca gcaacatttc tctcatagn 329

<210> 2772
<211> 302
<212> nucleic acid
<213> Glycine max
<400> 2772

tgncgcntac ntaagnncgg aattcggctc gnggcgaaga agaagccaca gagaaccagt 60
ctcctactct ctctaccca caagaaaaat ggcattctac atgcgttgga tcccccgca 120
tggttaccga agagagagct caagtctcgt ctcgagtctt tctgggatgg caagagcagc 180
gccgaggatt tgcagaaggt ggctgctgat ctcaggtcat ccatnctgga agcagatggc 240
tggtgctggg atcaagtaca tcccnagcaa cacttctcgn tctatgacca gctgcnnacg 300
cc 302

<210> 2773
<211> 281
<212> nucleic acid
<213> Glycine max
<400> 2773

gcacgcgtac gtaagctcgg aattcggctc ganctcgagc cgattcggct cgagttgctc 60
cctcagaagc gaagaaacng ccacagagaa ccagtctcct actctctctc aaccacaaga 120
aaaatggcat ctcacatcgt tggatacccc cgcattgggtc ccaaggagag ctcaagttcg 180
ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtggctgctg 240
atctcaggtc atccatctgg aagcagatgg ctgggtgctgg g 281

<210> 2774
<211> 286
<212> nucleic acid
<213> Glycine max
<400> 2774

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctgcttca acttgctccc 60
tcagaagcga agaagaagcc acagagaact agtctcctac tctcaccgcg aagaaaaaaa 120
tggcatctca catcgttgga taccctcgta tgggtcccaa gagagagnnc aagttcgctc 180
tcgagtcttt ctgggatggc aagagcagcg ccgaggattt gcagaagggtg tcttctgac 240
tcagggcacc atctggaagc agatggctga tgctgggac aagtac 286

<210> 2775
<211> 310
<212> nucleic acid
<213> Glycine max

<400> 2775

cncngcacnc gtacgtaagc tcggaattcg gctcgagctt gtcctctcag angcgaagaa 60
gaagccacag agaantagtc tctactctc acccgccaag anaaaaatgg cattctcaca 120
tcogttggat accctcgat ggggtcccaag agagagctca agttcgctct cgagtcttct 180
gggatggcaa gnncagcgcg gaggatttgc agaagggtgc tctgatctca gggcatccat 240
ctggaagcag atggctgatg ctgggatcaa gtacatcccc agcaacattt ctctcatatg 300
accaggttct 310

<210> 2776
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 2776

tngcangcac gcgtacgtaa gctcngaatt cngcnccgag ctctgcttca acttgctccc 60
tcagaancga agaagaagcn acagagaact agtctcctac tctccaccgc caagaaaaaa 120
atggcatctc acatcgttgg ataccctcg atgggtccca agagagagct acaagttcgc 180
tctcgagtct ttctgggatg gcaagagcag cgcgaggat ttgcagaagg tgtcttctga 240
nctnagggca tccatctgga agcagatggc nnatgctggg atcattacat cccagcaa 299

<210> 2777
<211> 253
<212> nucleic acid
<213> Glycine max

<400> 2777

cttgctcnnt naganncgaa gaagaagcca cagaggacta gtctnctac tgctcaccgc 60
caagaaaaaa atggcatctc anacccgttg gatacccccg tatgggtcnc aanagagagc 120
tcaagttngc tgctcgagtc tttctgggat ggcaatagca gngccganga tttgcagaag 180
gtgtcttctg atctcagggc atccatctgg aatcagatgg ctgatgcngg gatcaagtnc 240
atccccagca aca 253

<210> 2778

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2778

gcaaangcac gcgagacgta anctcggaat tcggctcgag naacttgntc cctcagnagc 60
gaagaagaag cnacananaa ctagtctcct acnncnaacc gnaagaaaaa natggcatct 120
cacatcggtg gatacccccg tatnggtngc aanagagagc tcaagttcgc tctcgagtct 180
tnctgngatg gnaagannag cgccgaggat ttgcagaagg tgtcttctga tctcagggca 240
tccatctgga agcagatggc tangctggga tcaagtacat cccangca 288

<210> 2779

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2779

aaatnctana agtcgcangc acgcgnacgt aanctcgga ntcggctcga gnaacttgct 60
ccctcagaag cgaagaagaa gccacanaga actagtctcc tactgctcac ccgcnagaaa 120
aaaatggcat ctncacatnc gttggatanc cccgnatgng ngcccaagan agagctcaag 180
ttcgctctcg agtcttctctg ggatggcagn agcagcgccg aggatttgca gaagggtgtct 240
tcngantca gggcatccat ctngaagcag atngctgatg ctgggatcaa gnacatctcc 300
aggaacactt tctctnactn 320

<210> 2780

<211> 249

<212> nucleic acid

<213> Glycine max

<400> 2780

ctgcntcaac ttgctccctg caganncgaa ncaagaagcc acagagnact agtctnccta 60
ctctcaccgc caananaaaa atggcatctc acatncgttg gatancccg tatgggtccc 120
aaganagagn tcaagttcgc tctcgagtct ttctgggatg gnaagagcag cgccgaggat 180
ttgcagaagg tgtnttctga tctcagggna tncatctgga agnagatggc tgatgntggg 240
ntcaagtac 249

<210> 2781

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2781

acggcacang nacgtaant cggaattcgg ctcgagcann cttgctccct cagaagcgaa 60
gaagaagcca cagagaacta gtctcctact notcaccgc aagaaaaaaa tggcatctca 120
catacgttgg atacccccgn atgggtccca agagagagct caagtctcgt ctcgagtctt 180
tctgggatgg caagagcagc gccgaggatt tgcagaaggt gtctcngatc tcagggcatc 240
catctggaag cagatggctg atgctgggat nagtacann ccagcaacat ttctctcata 300

<210> 2782

<211> 262

<212> nucleic acid

<213> Glycine max

<400> 2782

gtcgcaangca cgcgtaacgt naagctncgg taattcggcn cgaggccaca gagaactagt 60
ctncctactc tcnnccgcaa gaaaaaatg gcatctcaca tcgttgata ccccgatatg 120
gggcccaaga gagagctcaa gttcgctctc gantctttct gggatggcaa gagcagcgcc 180
gaggatttgc agaaggtgtc tctgatctca gggcatccat ctggaagcag atgctgatgt 240
ggtcaagnac tcccgcacan tt 262

<210> 2783

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 2783

tgcgatgcac gcgtacgtaa gtcggaatt cggctcgagc aacttgctcc ctcagaagcg 60
aagaagaagc cacagagaac cagtctccta ctctctctac acccacaaga aaaatggcat 120
ctcacatcgt tggatacccc cgcattgggtc ccaagagaga gctcaagttc gctctcgagt 180
ctttctggga tggcaagagc agcgccgagg atttgcagaa ggtggctgct gatctcaggt 240
ca 242

<210> 2784

<211> 308

<212> nucleic acid

<213> Glycine max

<400> 2784

ncgcntgcac gcgtacgtaa gtcggaatt cggctcgaga acttgctccc tcagaagcga 60
agaagaagcc acagagaact agtctcctac tcttcacccg caagaaaaaa atggcatctc 120
acatcgttgg atacccccgt atgggtccca agagagagct caagtctcgt ctcgagtctt 180
tctgggatgg caagagcagc gccgaggatt tgcagaaggt gtcttctgat ctcagggcat 240
ccatctgnan canatggctg atnncgggnt ncagtacatc cccagcaaca tttcttctca 300
tatgacca 308

<210> 2785

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2785

cttgctccct ccagaagacg aagaagaagc cacagagaat agttctccta ctctncaacc 60
gnaaganaaa acatggcatc tcacatcgtt ggatacccc gtatgggtcc caagagagag 120
ctcaagttcg ctctcgagtc tttctgggat ggcaagagca gngccgagga tttgcagaag 180
gtgtctctga tctcagggca tcccatctgg naggagatgg ctgatgtggg atnngtacat 240
cccagcaaca tttctcncat cngacangtt ctcgacg 277

<210> 2786

<211> 280
 <212> nucleic acid
 <213> Glycine max

 <400> 2786

 tatnaacttg ctcnntcaga ncgcgaaagaa gaagnacacag agaanchagt ctccctantnc 60
 tcaccccgnaa gnnaaaaatg ggcattctcn catcggttga taacccccgt atgggtgcc 120
 agagagagct caagttcgnt ctncagtcct tctgggatgg caagagcagc gncngaggat 180
 ttgcagaagg tgtcttctga tctcaggnc tccatctgga agcagatggc tgntgntggg 240
 atcagtacat cccnagcaac acttctctca ctatgaccag 280

<210> 2787
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2787

 gcacncgtac gtaagctcgg aattcggctc gagcttgctc cctcanaagc gaagnagaag 60
 ccacananaa ccagtctcct actctctctc acccacaaga naaatgggat ctcacatcgt 120
 tggntacccc cgcattgggtt cccaaganag agctcaagtt cgtctctcag tctttctggg 180
 ntggcnagag cagcgccgag gatttgcaga aggtggctgc tgatctcagg tcatccaatc 240
 tggaancaag attgccngat cggggatcaa gctccatncc cagcaacann tttttgct 298

<210> 2788
 <211> 151
 <212> nucleic acid
 <213> Glycine max

 <400> 2788

 gtgcangca ntgtacgtaa gctcggaatt cggctcgaga tctggaagca gatggctgat 60
 gctgggatca agtacatccc cagcaacact ttctctcact atgaccaggt tctcgacgcc 120
 accgccaccc tcggtgccgt tccaccaagg t 151

<210> 2789
 <211> 234
 <212> nucleic acid
 <213> Glycine max

<400> 2789

gcacgcgtac gtaagctcgg aattcggctc gagcttcaac ttgctccctc agaagcgaag 60
aagaagccac agagnactag tctctactct tcaccogcaa naaaaaaatg ggcatctcac 120
attcgttgga tcccccgta tgggtcccaa gagagagctc aagttcgctc tcgagtcttc 180
naggatggca agagcagcgc cgaggatttg cagaaggtgt cttctgatct cagg 234

<210> 2790

<211> 138

<212> nucleic acid

<213> Glycine max

<400> 2790

gnagaagaag ccacagagaa ctagtctcct actctcacc cgaagaaaaa aatggcatct 60
cacatcnttg gatacccccg tatgggtccc aagagagagc tcaagttcgc tctcgagtct 120
ttctgggatg gcaagagt 138

<210> 2791

<211> 152

<212> nucleic acid

<213> Glycine max

<400> 2791

acacgcgtac gtaaagctnc ggaattcngc tcgagctgga agcagatggc tgatgctggg 60
atcaagtaca tccccagcaa cactttctnn tnactatgac caggttctcg acgccacccg 120
ccaccctcgg tgccgttcca cncangnnag gn 152

<210> 2792

<211> 501

<212> nucleic acid

<213> Glycine max

<400> 2792

gtttgatctc cgtggnccta tataaagaga tatgangtcg nntccacgcg tangtaanac 60
tcggaattcg gctcgagcaa onttccaatc cttccaacca ccactantgg atccttonct 120
canacngtag aactgaggag ggtacgccgt gagttcaagg ctaacatgat ctccgaggaa 180
gagtatgtta agtcaattaa ggaggaaatt cncaaagttg ttgaacttca anaagagctt 240

gatattgatg ttcttgttca tggagaacca gagagaaatg atatggttga gtacntcggt 300
gagcaattgt caggctttgc cttcactgtt aatgggtggg tgcaatccta tggttcccgt 360
tgtgtgaagc cacnaatcat ctatggtgat gtgagccgcc caaagccaat gactgtcttc 420
tggtcancctc tggctcagan ctttaccag cgccnaatga agggaatgct taccgggtccg 480
gttaccaatc ccaactgggn c 501

<210> 2793
<211> 412
<212> nucleic acid
<213> Glycine max
<400> 2793

gagaagacga cagaaggggg ctaaggcatt gtctggcaac aaggatgtgg ctttcttctc 60
tgctaattgt gcanctcang cttcaaggaa gtccctctcca anagtgacca acnaggctgt 120
tcagaaggct gctgctgcat tgaagggttc agatcatcnc cgtgcaacaa atgtcagtgc 180
cagactggat gctcaacaaa anaagctcaa ccttccaatc cttccaacca ccactattgg 240
atccttccct cagactgtan aactgaggag ggtacgcogt gagttcaagg ctaacaagat 300
ctccgaggaa gagtatgtta agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca 360
agaagagctt gatattgatg ttcttgttca tgganaacca anaganaaat ta 412

<210> 2794
<211> 350
<212> nucleic acid
<213> Glycine max
<400> 2794

cacactgctg ttgatcttgt taacgagacc aagttggatg acgagatcaa gtcattggcta 60
gcatttgctg cacaaaaaat tgttgaagtt aacgcattgg ctaaggcatt gtctggcaac 120
aaggatgtgg ctttcttctc tgctaattgt gcagctcagg cttcaaggaa gtccctctcca 180
agagtgacca acgaggctgt tcagaaggct gctgctgcat tgaagggttc agatcatcgn 240
cgtgcaacaa atgtcagtgc cagactggat gctcaacaaa agaagctcaa ccttccatcc 300
ttccaaccac cactattgga tccttccctc agactgtaga actgaggagg 350

<210> 2795

<211> 454
 <212> nucleic acid
 <213> Glycine max

<400> 2795

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ccacgcgtcc ggatcatcgc cgtgcaacaa atgtcagtgc cagactggat gctcaacaaa 60
agaagctcaa ngttccngtc cttccaacca ccactattgg atccttccct cagactgtag 120
aactgangag ggtacgccgt gagttcaagg ctaacaagat ctccgaggaa gagtatgtta 180
agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca anaagagctt gatattgatg 240
ttcttgttca tggagaacca gagagaaatg atntgggtga gtacttcggt gagcaattgt 300
caggctttgc cttcactggt aatgggtggg tgcaatctat ggggccggt gtgtgaaagc 360
caccaattca tctatgggtg aatgtgnaaa nncngtccaa aagccaatga ctgtcttctg 420
gtcatctntg gcttaaangc tttaccaaag cgct 454
```

<210> 2796
 <211> 446
 <212> nucleic acid
 <213> Glycine max

<400> 2796

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cgaggctggt cagaaggctg ctgctgcatt gaagggttcn gatcatcgcc gtgcaacaaa 60
tgtcagtgcc agactggatt ctcaacaaaa gaagctcaac cttccaatcc tgccaaccac 120
cactattgga tcttccctc agactgtaga actgaggagg gtacgccgtg aattcaaggc 180
taacaagatc tccgaggaag agtatgtaaa gtcaattaag gaggaaattc gcaaagttgt 240
tgagcttcaa gaagagcttg atattgatgt tcttgggtcat ggagaaccag agagaaatga 300
tatgggtgag tactttcggg gaacaattgt caagctttgg cntnaccggt aatgggtngg 360
tgcaatccta tggttcccggt tcgtgaaanc cccgatcatt tatgggaatg ttagccgccc 420
aaagccatga ccgntttttg gtattt 446
```

<210> 2797
 <211> 489
 <212> nucleic acid
 <213> Glycine max

<400> 2797

ttctctgccc gttcngtach nntctccnaa ttcgngggcc gacccacgcn tccggcagat 60
 ctgagaccac ctaccagatt gctttgtcta tcaaggacga agtggaagac cttgaaaagg 120
 ctggcatcac tggtatccaa attgatgaag ctgctttgag agaggggtctt ccaactgagga 180
 aatcagagca agctcactac ttggactggg ctgtccatgc cttcagaatc accaatgttg 240
 gtgtccagga taccacccag gtacactctt ttggatcatc gcaaatact gaattagaaa 300
 ttttttttgt tcattctcat tttcacatat gttgtaataa tcaacttttc gtattgacag 360
 atccacactc acatgtgcta ctggaacttc aacgacatca tccactccat catcgacatg 420
 gacgccgatg ttatcaccat tgagaaaatc cgccccgacg anaancttcc gtcagtcttc 480
 cgccaangg 489

<210> 2798
 <211> 340
 <212> nucleic acid
 <213> Glycine max
 <400> 2798

tgcgancnc gcgtacgtaa gctcggaatt cggtctgagg ntcaagtcac ggctagcttt 60
 tgctgccccaa aaaattgttg aagttaacgc attggctaaa gcattgtctg gccacaagga 120
 tgangccttc ttctctggta atgctgctgc tctggcttca aggaagtctt ctccaagagt 180
 gaccaacgag gctgttcaga aggctgctgc tgcattgaag ggttcagatc atcgccgtgc 240
 aanaaatgtc agtgccagac tggattctca acaaaagaag ctcaaccttc caatcctgcc 300
 aaccaccact attggatcct tccctcagac tgtagaactg 340

<210> 2799
 <211> 317
 <212> nucleic acid
 <213> Glycine max
 <400> 2799

gtcgancgc agcgtacgta agctcggaat tcggctgag aaatgatatg gttgagtact 60
 tcggtgagca attgtcaggc tttgccttca ctgttaatgg gtgggtgcaa tcctatgggt 120
 cccgttgtgt gaagccacca atcatctatg gtgatgtgag ccgccccaaag ccaatgactg 180
 tcttctggtc atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg 240

gtcctgttac cattctcaac tggtcctttg ttagaaatga ccaacctaga tctgagacca 300
cctaccagat tgctttg 317

<210> 2800
<211> 317
<212> nucleic acid
<213> Glycine max
<400> 2800

gtcgngcac gcgtacgtaa gctcggaatt cggtcgcaga cgaagtggag gaccttgaaa 60
aggctggcat cactgttatc caaattgatg aagctgcttt gagagagggg ctgccactga 120
ggaaatcaga acaagctcac tacttggact gggctgtcca tgccttcaga atcaccaatg 180
ttgggtgtgca ggataccact cagatccaca cccacatgtg ctactccaac ttcaacgaca 240
tcatccactc catcatcgac atggacgctg atgttatcac cattgagaac tctcgctccg 300
atgagaagct cctgtca 317

<210> 2801
<211> 337
<212> nucleic acid
<213> Glycine max
<400> 2801

cnngtcgan gcacgcntac gtaagctcgg aattcggctc gaggccacca atcatctatg 60
gtgatgtgag ccgccc aaag ccaatgactg tcttctggtc atctctggct cagagcttta 120
ccaagcnccc aatgaaggga atgcttaccg gtctgttac cattctcaac tggtcctttg 180
ttagaaatga ccaacctaga tctgagacca cctaccagat tgctttggct atcaaggacg 240
aagtggagga ccttgaaaag gctggcatca ctgttatcca aattgatgaa gctgctttga 300
gagagggctc gccactgagg aaatcagaac aagctcn 337

<210> 2802
<211> 329
<212> nucleic acid
<213> Glycine max
<400> 2802

cangcacgcg tacgtaagct cggaattcgg ctcgagntgc tgcattgaag gggtcagatc 60

atcgccgtgc aacaaatgtc agtgccagac tggatgctca acaaaagaag ctcaaccttc 120
 caatccttcc aaccaccact attggatcct tccctcagac tgtagaactg aggagggtag 180
 gccgtgagtt caaggctaac aagatctccg aggaagagta tgттаagtca attaaggagg 240
 aaattcgcaa agttgttgaa cttcaagaag agcttgatat tgatgttctt gttcatggag 300
 aaccagagag aatgatatg gttgagtac 329

<210> 2803
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 2803

tcgcangcac gcgtacgtaa gctcggaatt cggtcgagc tgaggaaatc agagcaagct 60
 cactacttgg actgggctgt ccatgccttc agaataacca atgttggtgt ccaggatacc 120
 acccagatcc aactcacat gtgctaactg aacttcaacg acatcatcca ctccatcatc 180
 gacatggacg ccgatgttat caccattgag aactctcgct ccgacgagaa gcttctgtca 240
 gtcttccgcg aaggtgtgaa gtatggtgct ggaattggcc ctggtgtcta tgacatccac 300
 tccccaaaga tacc 314

<210> 2804
 <211> 328
 <212> nucleic acid
 <213> Glycine max
 <400> 2804

ctcangcacg cntacgtaan ctcggaattc ggctcgagng tacgccgtga attcaaggct 60
 aacaagatct ccgaggaaga gtatgtaaag tcaattaagg aggaaattcg caaagttggt 120
 gagcttcaag aagagcttga tattgatgtt cttgttcatg gagaaccaga gagaaatgat 180
 atggttgagt acttcggtga acaattgtca ggctttgcct tcaccgttaa tgggtgggtg 240
 caatcctatg gttcccggtg cgtgaagcca ccgatcatct atggtgatgt gagccgcca 300
 aagccaatga ccgtcttctg gtcattctc 328

<210> 2805
 <211> 323
 <212> nucleic acid

<213> Glycine max

<400> 2805

ngtcgcangc acgcgtacgt aagctcggaa ttccggctcga gggcattgtc tggcaacaag 60
gatgtggcct tcttctctgc taatgctgca gctcaggcct caaggaagtc ctctccaaga 120
gtgaccaacg aggcgtgtca gaaggctgct gctgcattga agggttcaga tcatcgccgt 180
gcaacaaatg tcagtgccag actggatgct caacaaaaga agctcaacct tccaatcctt 240
ccaaccacca ctattggatc cttccctcag actntagaac tgaggagggt acgccgtgag 300
ttcaaggcta acaagatctc cga 323

<210> 2806

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2806

ngnncgcatg cacgcgtacg tnaagctcga attcggctcg agatatgggtt gagtacttcg 60
gtgagcaatt gtnaggcttt gccttcaactg ttaatgggtg ggtgcnatcc tatggttccc 120
gttgtgtgaa gccaccaatc atctatgggtg atgtgagccg cccaaagcca atgactgtct 180
tctggtcacg tctggctcag agctttacca agcgcccaat gaagggaatg cttaccggtc 240
ctgttaccat tctcaactgg tcctttgtta ganatgacca acctagatct gagaccacct 300
accagattgc tt 312

<210> 2807

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 2807

canngacgcg tacgtnagct cggaattcgg ctccaggaaa tgatatgggtt gagtacttcg 60
gtgagcaatt gtcaggcttt gccttcaactg ttaatgggtg ggtgcaatcc tatggttccc 120
gttgtgtgaa gccaccaatc atctatgggtg atgtgagccg cccaaagcca atgactgtct 180
tctggtcacg tctggctcag agctttacca agcgcccaat gaagggaatg cttaccggtc 240
ctgttaccat tctcaactgg tcctttgtta gaaatgacca acctagatct gagac 295

gaaatgatat ggttgagtac ttcggtgagc aattgtcagg ctttgccctc actgttaatg 240
 ggtgggtgca atcctatggt tcccgttgtg tgaagccacc aatcatctat ggtgatgtga 300
 gccgcc 306

<210> 2811
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2811

acnagcgang cacgcgtacg taagctcgga attcggctcg agctccatca tcgacatgga 60
 cgctgatgtt atcaccattg agaactctcg ctccgatgag aagctcctgt cagtcttccg 120
 tgaaggtgtg aagtatggtg ctggaattgg cctggtgtc tatgacatcc actccccaag 180
 aataccacca actgaagaaa tcgctgacag aatcaataag atgcttgacg tgctcgagaa 240
 gaacatcttg tgggtcaacc ctgactgtgg totcaagacc cgcaagtaca ctgaagtga 300
 gccagccctc 310

<210> 2812
 <211> 353
 <212> nucleic acid
 <213> Glycine max
 <400> 2812

nnnnnnaaaa gtcgggaagt cgcangcacg cgtacgtaag ctcggaattc ggctcgnanc 60
 tcgagccgaa tcggctcgag ccagattgct ttggctatca aggaagnagt ggaggacctt 120
 acnaaggctg gcatcactgt actccaaatt gatgaagctg ctttgagaga gggctctgcca 180
 ctgaggaaat cagaacaagc tcactacttg gactgggctg tccatgcctt cagaatcacc 240
 aatgttggtg tgcaggatac cactcagatc cacaccacac tgtgctactc caacttcaac 300
 gacatcatcc actccatcat cgacatggac gctgatgtta tcaccattga gat 353

<210> 2813
 <211> 297
 <212> nucleic acid
 <213> Glycine max
 <400> 2813

ctcnncgcgt tacgtaagct cggaattcg gctcgaggct gctttgagag agggctctgcc 60
actgaggaaa tcagaacaag ctactactt ggactgggct gtccatgcct tcagaatcac 120
caatgttggt gtgcaggata ccactcagat ccacaccac atgtgctact ccaacttcaa 180
cgacatcatc cactccatca tcgacatgga cgctgatgtt atcaccattg agaactctcg 240
ctccgatgag aagctcctgt cagtcttccg tgaagggtgt aagtatgggt ctggaat 297

<210> 2814
<211> 551
<212> nucleic acid
<213> Glycine max
<400> 2814

gnnngagagg tttntgntan gggggaaggg ggggnanatn ntagaanngc tatgacgtcg 60
catgcacgcg tacgtaagct cggaattcgg ctcgagtgga cgctgatgtt atcaccattg 120
agaactctcg ctccgatgag aagctcctgg tcagtcttcc gtgaagggtgt gaagtatgggt 180
gctggaattg gccctggtgt ctatgacatc cactccccaa gaataccacc aactgaagaa 240
atcgctgaca gaatcaataa gatgcttgca gtgctcgaga agaacatctt gtgggtcaac 300
cctgactgtg gtctcaagac ccgcaagtac actgaagtga agccagccct cacaacatg 360
gttgccgcag caaaactcat ccgtaacgaa cttgccaagt gaatgggtata aagaaagtag 420
aatctacaag ttcatgtgtt ctgcttttat tataccncca aggaaaaatt ttctatantn 480
gggtggttca aataaccggt gtggaatatt tanaggttta acatgctctg tgagcaattg 540
atctttctca c 551

<210> 2815
<211> 336
<212> nucleic acid
<213> Glycine max
<400> 2815

agtntcann nntcanacgt cgcacgcac cgtacgtaag ctcggaattc ggctcgagga 60
tcacgcgcgt gcaacaaatg tcagtgccag actggattct caacaaaaga agctcaacct 120
tccaatctcg ccaaccacca ctattggatc cttccctcag actgtagaac tgaggaggggt 180
acgccgtgaa ttcaaggcta acaagatctc cgaggaagag tatgtaaagt caattaagga 240

ggaaattcgc aaagttgttg agcttcaaga agagcttgat attgatgttc ttgttcacgg 300
agaaccagag agaaatgata tggttgagta cttcgg 336

<210> 2816
<211> 313
<212> nucleic acid
<213> Glycine max

<400> 2816

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gactcgaact tcaacgacat 60
catccactcc atcatcgaca tggacgccga tgttatcacc attgagaact ctgctccga 120
cgagaagctt ctgtcagtct tccgcgaagg tgtgaagtat ggtgctggaa ttggccctgg 180
tgtctatgac atccactccc caagaatacc accaactgaa gaaattgctg acagaatcaa 240
caagatgctg gcagtgtctg agangaacat cttgtgggnt gaacctgact gtgggctcaa 300
gacccgtaaa gtn 313

<210> 2817
<211> 304
<212> nucleic acid
<213> Glycine max

<400> 2817

gtcgcangca cgcgtacgta agctcggaa tggctcgag ngctgctttg agagagggtc 60
ttccactgag gaaatcagag caagctcact acttggactg ggctgtccat gccttcagaa 120
tcaccaatgt tgggtgtccag gataccaccc agatccacac tcacatgtgc tactcgaact 180
tcaacgacat catccactcc atcatcgaca tggacgccga tgttatcacc attgagaact 240
ctgctccga cgagaagctt ctgtcagtct tccgcgaagg tgtgaagtat ggtgctggaa 300
ttgg 304

<210> 2818
<211> 438
<212> nucleic acid
<213> Glycine max

<400> 2818

ccacgcgtcc gaagagagaa atgatatggn tgngtttttn ngagaacaat ngnnangctt 60

<210> 2821
 <211> 323
 <212> nucleic acid
 <213> Glycine max

<400> 2821

tcncatgcac gcgtacgtaa gctcgggaatt cggctcgagc tagcttttgc tgcccaaaaa 60
 attgttgaag ttaacgcatt ganctaaagc attgtctggc cacaaggatg aggccttctt 120
 ctctggtaat gctgctgctc tggcttcaag gaagtcttct ccaagagtga ccaacgaggc 180
 tgttcagaag gctgctgctg cattgaaggg ttcagatcat cgccgtgcaa caaatgtcag 240
 tgccagactg gattctcaac aaaagaagct caaccttcca atcctgccaa ccaccactat 300
 tggatccttc cctcagactg tag 323

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<210> 2822
 <211> 290
 <212> nucleic acid
 <213> Glycine max

<400> 2822

acgtnagctc ggaattcggc tcgagcttgg actgggctgt ccatgccttc agaatcacca 60
 atgttggtgn gcaggatacc actcagatcc acaccacat gtgctactcc aacttcaacg 120
 acatcatcca ctccatcatc gacatggacg ctgatgttat caccattgag aactctcgct 180
 ccgatgagaa gtcctgtca gtcttccgtg aagggtgtgaa gtatgggtgct ggaattggcc 240
 ctggtgtcta gacatccact cccaagaat accaccaact gaagaaatcg 290

<210> 2823
 <211> 265
 <212> nucleic acid
 <213> Glycine max

<400> 2823

ctgctgttga tcttgtaac gagaccaagt tggatgacga gatcaagtca tggctagcat 60
 ttgctgcaca aaaaattggt gaagttaacg cattggctaa ggcattgtct ggcaacaagg 120
 atgtggcctt cttctctgct aatgctgcag ctgaggcttc aaggaagtcc tctccaagag 180
 tgaccaacga ggctgttcag aaggctgctg ctgcattgaa gggttcagat catcgccgtg 240

caacaaatgt cagtgccaga ctgga

265

<210> 2824
<211> 289
<212> nucleic acid
<213> Glycine max

<400> 2824

cgtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gcaaagttgt tgaacttcaa 60
gaagagcttg atattgatgt tcttgttcat ggagaaccag agagaaatga tatggttgag 120
tacttcggtg agcaattgtc aggccttggc ttcactgtta atgggtgggt gcaatcctat 180
ggttcccggt gtgtgaagcc accaatcatc tatggtgatg tgagccgccc aaagccaatg 240
actgtcttct ggtcatctct ggctcagagc tttaccaagc gcccaatgn 289

CGTCG
CATGC
ACGCG
TACGT
AAGCT
CGGAA
TTCGG
CTCGA
GCAAG
TTGT
TGAAC
TTCAA
GAAGA
GCTTG
ATATT
GATGT
TCTTG
TTCAT
GGAGA
ACCAG
AGAGA
AATGA
TATGG
TTGAG
TACTT
CGGTG
AGCAA
TTGTC
AGGCT
TTGGC
TTCAC
TGTTA
ATGGG
TGGGT
GCAAT
CCTAT
GGTT
CCCGT
GTGTG
AAGCC
ACCA
ATCAT
C
TATGG
TGATG
TGAGC
CGCCC
AAAGC
CCAAT
G

<210> 2825
<211> 265
<212> nucleic acid
<213> Glycine max

<400> 2825

gttaacgcat tggctaaggc attgtctggc aacaaggatg tggccttctt ctctgctaata 60
gctgcagctc aggccttcaag gaagtcctct ccaagagtga ccaacgaggc tgttcagaag 120
gctgctgctg cattgaaggg ttcagatcat cgccgtgcaa caaatgtcag tgccagactg 180
gatgctcaac aaaagaagct caaccttcca atccttccaa ccaccactat tggntccttc 240
cctcagactg tagaactgag gaggg 265

<210> 2826
<211> 304
<212> nucleic acid
<213> Glycine max

<400> 2826

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gcttcacact gctgttgatc 60
nagttaacga gaccaagttg gatgatgaga tcaagtcatg gctagctttt gctgccccaa 120
aaattgttga agttaacgca ttggctaaag cattgtctgg ccacaaggat gaggccttct 180
tctctggtaa tgctgctgct ctggcttcaa ggaagtcttc tccaagagtg accaacgagg 240

ctgttcagaa ggctgctgct gcattgaagg gttcagatca tcgccgtgca acaaattgtca 300
gtgc 304

<210> 2827
<211> 269
<212> nucleic acid
<213> Glycine max

<400> 2827

ctggtttgan tgttcttggt gagacctact ttgctgacat ccctgctgag gcatacaaga 60
ccctcacatc tctgaatggc gtcactgcat atggatttga ttggtccgt ggaaccaaca 120
ctcttgattt gatcaagggg ggatttccca gcggaaaata cctctttgct ggagtgggtg 180
atggaaggna catctgggcc aatgaccttg ctgcttctct cactaccttg cagggtcttg 240
agggcattgt gggcaaagat aagcttggt 269

<210> 2828
<211> 315
<212> nucleic acid
<213> Glycine max

<400> 2828

tcnatnnag cgnaggtann tacgtaagct cggaattcgg ctcgagtacg gctgcgagaa 60
gacgacagaa gggcatggag aaccagagag caatgatatg gttgagtact tcggtgagca 120
attgtcaggc ttgctcttca ctgttaatgg gtgggtgcaa tcctatgggt cccgttgtgt 180
gaagccacca atcatctatg gtgatgtgag ccgccccaaag ccaatgactg tcttctggtc 240
atctctggct cagagcttta ccaagcggc aatgaaggga atgottaccg gtctgtttac 300
cattctcaac tggtc 315

<210> 2829
<211> 320
<212> nucleic acid
<213> Glycine max

<400> 2829

gtcgcangca cgcgtacgtn agctcggnat tcggctcgag caagcgccca atgaaggga 60
tgcttaccgg tctgtttacc attctcaact ggtcctttgt tagaaatgac caacctagat 120

<210> 2832
 <211> 323
 <212> nucleic acid
 <213> Glycine max

 <400> 2832

 nnaagcntc angnacgcgt acgttagctc ggaattcggc tcgagggcctt caaggaagtc 60
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 tntcgccgt gcaacaaatg tcagtgccag actggatgct caacaaaaga agctcaacct 180
 tccaatcctt ccaaccacca ctattggntc cttccctcag actgtagaac tgaggaggga 240
 ccgccgtgag ttcaaggcta acaagatctc cgaggaagag tatgttaagt caattaagga 300
 ggaaattcgc aaagttgttg aan 323

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<210> 2833
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 2833

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 cgcaaagttg ttgaacttca agaagagctt gatattgatg ttcttggttca tggagaacca 120
 gagagaaatg atatggttga gtacttcggt gagcaattgt caggctttgc cttcactggt 180
 aatgggtggg tgcaatccta tggttcccgt tgtgtgaagc caccaatcat ctatggtgat 240
 gtgagccgcc caaagccaat gactgtcttc tggctcatctc tggctcagag ctttac 296

<210> 2834
 <211> 265
 <212> nucleic acid
 <213> Glycine max

 <400> 2834

 cgcccaatga agggaatgct taccggtcct gttaccattc tcaactggtc ctttggttaga 60
 aatgaccaac ctagatctga gaccacctac cagattgctt tgtctatcaa ggacgaagtg 120
 gaagaccttg aaaaggctgg catcactggt atccaaattg atgaagctgc tttgagagag 180
 ggtcttcac tgaggaaatc agagcaagct cactacttgg actgggctgt ccatgccttc 240

agaatcacca atgttggtgt ccagg

265

<210> 2835
<211> 325
<212> nucleic acid
<213> Glycine max

<400> 2835

wgtngcnngc acgcgtacgt aagctcggaa ttccggctcga gttcccagtg gaaaataacct 60

ctttgctgga gtggttgatg gaaggaacat ctgggccaat gaccttgctg cttctctcac 120

tacattgcag ggtcttgagg gcattgtggg caaagataag cttgttgtgt ccacctctc 180

ctcccttctt cacactgctg ttgatcttgt taacgagacc aagttggatg acgagatcaa 240

gtcatggcta gcatttgctg cacaaaaaat tgttgaagtt aacgcattgg ctaaggcatt 300

gtctggcaac aaggagtggc cttct 325

<210> 2836
<211> 346
<212> nucleic acid
<213> Glycine max

<400> 2836

gttaacgcat tggctaaggc attgtctggc aacaaggatg tggccttctt ctctgctaatt 60

gctgcagctc aggcttcaag naagtcctct ccnagagtga ccaacgaggc tgttcagaag 120

gctgctgctg cattgaaggg ttcanatcat cgccgtgcaa caaatgtcag tgccagactg 180

gatgctcaac aaaagaagct caaccttcca atccttccaa ccaccactnt tgnntccttc 240

cctcagactg tagaactgag gagggtagc gtgagttcaa ggtaacaaga ntccgaggaa 300

gagtatgtta agnccattaa ggaganatnt caagtgtgaa ctnaag 346

<210> 2837
<211> 312
<212> nucleic acid
<213> Glycine max

<400> 2837

tcgcangcac gcgtacgtaa gctcgggaatt cggctcgagg accttgctgc ttctctcact 60

acattgcagg gtcttgaggg cattgtgggc aaagataagc ttgttgtgtc cacctcctcc 120

tcccttcttc acactgctgt tgatcttggt aacgagacca agttggatga cgagatcaag 180
 tcatggctag catttgctgc acaaaaaatt gttgaagtta acgcattggc taaggcattg 240
 tctggcaaca aggatgtggc cttcttctct gctaattgctg cagctcaggc ttcaaggaag 300
 tcctctccaa ga 312

<210> 2838
 <211> 307
 <212> nucleic acid
 <213> Glycine max
 <400> 2838

ntatccntnn acgtcgcang cacgcgtacg taagctcgga attcggtctg agngagggta 60
 cgccgtgagt tcaaggctaa caagatctcc gaggaagagt atgttaagtc aattaaggag 120
 gaaattcgca aagttgttga acttcaagaa gagcttgata ttgatgttct tgttcatgga 180
 gaaccagaga gaaatgatat ggttgagtac ttccgtgagc aattgtcagg ctttgccctc 240
 actgttaatg ggtgggtgca atcctatggt tcccgttggt tgaagccacc aatcatctat 300
 ggtgatg 307

<210> 2839
 <211> 310
 <212> nucleic acid
 <213> Glycine max
 <400> 2839

cttnnagtcn catncacnag tacgnaantc ngcnnnncng ttcttggtga gacctacttt 60
 gctgacatcc ctgctgaggc atacaagacc ctacatctc tgaatggcgt cactgcatat 120
 ggatttgatt tgggtccgtg aaccaacact cttgatttga tcaaggggtg atttcccagc 180
 ggaaaatacc tctttgctgg agtggttgat ggaaggaaca tctgggccaa tgaccttgct 240
 gcttctctca ctaccttgca ggttcttgag ggcattgtgg gcaaagataa gcttgttgtg 300
 tccacctcct 310

<210> 2840
 <211> 297
 <212> nucleic acid
 <213> Glycine max

<400> 2840

nttcgcatgc acgcgtacgt aagctcggaa ttccggctcga gcacctacca gattgctttg 60

tctatcaagg acgangtgga agaccttgaa aaggctggca tcaactgttat ccaaattgat 120

gaagctgctt tgagagaggg tcttccactg aggaaatcag agcaagctca ctacttggac 180

tgggctgtcc atgccttcag aatcaccaat gttgggtgtcc aggataccac ccagatccac 240

actcacatgt gctactcgaa cttcaacgac atcatccact ccatcatcga catggac 297

<210> 2841

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2841

gtcgtttnat gcacgcgtac gtaagctcgg aattcggctc gagaaaaatt gttgaagtta 60

acgcattggc taaggcattg tctggcaaca aggatgtggc cttcttctct gctaattgctg 120

cagctcaggc ttcaaggaag tcctctccaa gaggtagcaa cgaggctgtt cagaaggctg 180

ctgctgcatt gaagggttca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 240

ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattgga tccttccctc 300

aga 303

<210> 2842

<211> 241

<212> nucleic acid

<213> Glycine max

<400> 2842

gttcttggtc atggagaacc agagagaaat gatatgggtg agtacttcgg tgaacaattg 60

tcaggctttg cttcacctgt taatgggtgg gtgcaatcct atggttcccg ttgcgtgaag 120

ccaccgatca tctatggtga tgtgagccgc ccaaagccaa tgaccgtctt ctggtcattc 180

ctggctcaga gctttaccaa gcgcccaatg aagggaatgc ttaccgggtc tggttaccatt 240

c 241

<210> 2843

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2843

ntgcncgcta ngtaagctcg gaattcggct cgagtcaggc cttctttctct gctaagtctg 60
cagctcaggc ttcaaggaag tcctctccaa gaggtagcaa cgaggctggt cagaaggctg 120
ctgctgcatt gaaggggtca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 180
ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattggn tctttccctc 240
agactgtaga actgaggagg gtacgccgtg agttcaaggc tancaagatc tccgag 296

<210> 2844

<211> 265

<212> nucleic acid

<213> Glycine max

<400> 2844

cgcccaatga agggaatgct taccggctct gttaccattc tcaactggtc ctttgttaga 60
aatgaccaac ctagatctga gaccacctac cagattgntt tgtctatcaa ggacgaagtg 120
gaagaccttg aaaaggctgg catcactgtt atccaaattg atgaagctgc tttgagagag 180
ggctttccac tgaggaaatc agagcaagct cactacttgg actgggctgt ccatgccttc 240
agaatcacca angttgggtg ccagg 265

<210> 2845

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 2845

atntgtcgca tgcncgcta cgtaagctcg gnattcggct cgagngacat catccactcc 60
atcatcgaca ggncgccgat gttatcacca ttgagaantc tcgctccgac gagangcttc 120
tgtcagtctt ccgcgaaggt gtgangtatg gtgctggaat tggccctggt gtctatgaca 180
tccactcccc aagaatacca ccaactgaag anattgctga cagaatcaac aagatgctgg 240
cagtgtctga gaagnacatc ttgtgggtga nccctgactg tgggctcaag acccgtaagt 300
aactgaggt gaagc 315

<210> 2846

<211> 311
 <212> nucleic acid
 <213> Glycine max

 <400> 2846

 ngncatgcac gcgtacgtaa gctcgggaatt nggctcgagt cagatcatcg ccgtgcaaca 60
 aatgtcagtg ccagactgga ttctcaacaa aagaagctca accttccaat cctgccaaacc 120
 accactattg gatccttccc tcagactgta gaactgagga gggtagcgcg tgaattcaag 180
 gctaacaaga tctccgagga agagtatgta aagtcaatta angaggaant tcgcaaagtt 240
 gttgagcttc aagaagagct tgatatngat gttcttggtc atggagaacc agagagaaat 300
 gatatggttg n 311

<210> 2847
 <211> 256
 <212> nucleic acid
 <213> Glycine max

 <400> 2847

 gaagccaccg atcatctatg gtgatgtgag ccgccccaaag ccaatgaccg tcttctggtc 60
 atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg gtcctgttac 120
 cattctcaac tggtcctttg ttgaaatga ccaacctaga tctgagacca cctaccagat 180
 tgctttgtct atcaaggacg aagtggaaga ccttgaaaag gctggcatca ctgttatcca 240
 aattgatgaa gctgct 256

<210> 2848
 <211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 2848

 gtcgcatgca cgcgtacgta agctcggaat tcgggctcga ggatttgatc aagggtggat 60
 ttcccagcgg aaaatacctc tttgctggag tggttgatgg aaggaacatc tgggccaatg 120
 accttgctgc ttctctcact accttgacgg gtcttgaggg cattgtgggc aaagataagc 180
 ttgttggtgc cacctcctcc tcccttcttc aactgctgt tgatctagtt aacgagacca 240
 agttggatga tgagatcaag tcatggctag cttttgctgc caaaaaatt gttgaagtta 300

acgcatgg

308

<210> 2849
<211> 292
<212> nucleic acid
<213> Glycine max

<400> 2849

acaagnanna ngtacgtaag ctcggaatt cggtcgagc cttcttctct gctaattgctg 60
cagctcaggc ttcaaggaag tcctctccaa gaggtagcaa cgaggctggt cagaaggctg 120
ctgctgcatt gaaggggtca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 180
ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattgga tccttccttc 240
agactgtaga actgaggagg gtacgccgtg agttcaaggc taacaagatc tc 292

<210> 2850
<211> 292
<212> nucleic acid
<213> Glycine max

<400> 2850

cgccangcag cgtacgtaag ctcggaattc gggtcgagcc actgaggaaa tcgaacaagc 60
tcactacttg gactgggctg tccatgcctt cagaatcacc aatgttggtg tgcaggatac 120
cactcagatc cacaccacac tgtgtacttc caacttcaac gacatcatcc actccatcat 180
cgacatggac gctgatgtta tcaccattga gaactctcgc tccgatgaga agctcctgtc 240
agtcttccgt gaaggtgtga agtatgggtc tggaattggc cctgggtgtct at 292

<210> 2851
<211> 327
<212> nucleic acid
<213> Glycine max

<400> 2851

gtcncangca cgcntacgta anctcggaat tcgggtcgag aaatgatatg gttgagtact 60
tcgggtgaaca attgtcaggc ttgtccttca ccgttaatgg gtgggtgcaa tcctatgggt 120
cccgttgctg gaagccaccg atcatctatg gtgatgtgag ccgccccaaag ccaatnaccg 180
tcttctggtc atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg 240

gtcctgttac cattctcaac tggtcctttg ttagaaatga ccaacctang tataaactcc 300
acaccgaaaa atgaacatca aggaggg 327

<210> 2852
<211> 345
<212> nucleic acid
<213> Glycine max
<400> 2852

gtcgcangcn cgcgtacgtn agctcngaata tcggctcgag cnnagacct cacatctctg 60
aatnggcgtc actgcatatg gggttgattt ggcccggtga acccatactc ttgatttgat 120
caagggtgga tttcccagtg gaaaataacct ctttgctgga gtgggttgatg gaaggnacat 180
ctggggccaat gaccttgctg cttctctcac tacattgcag ggtcttgagg gcattgtggg 240
caaagataag cttgttggtg ccacctctc ctcctctctt cacactgtg ttgatcttgt 300
taacgagacc aagttggatg acgagatcaa gtcattggcta gcatt 345

<210> 2853
<211> 309
<212> nucleic acid
<213> Glycine max
<400> 2853

gtcgcnngca cgcgtacgta agctcggaaat tcggctcgag gctgttcaga aggctgctgc 60
tgcatgaag gggtcagatc atcgccgtgc aacaaatgac agtgccagac tggatgctca 120
acaaaagaag ctcaaccttc caatccttcc aaccaccact attggatcct tccctcagac 180
tgtagaactg aggagggtac gccgtgagtt caaggctaac aagatctccg aggaagagta 240
tgttaagtca attaaggagg aaattcgcaa agttgttgaa cttcaagaag agcttgatat 300
tgatgttct 309

<210> 2854
<211> 311
<212> nucleic acid
<213> Glycine max
<400> 2854

gtcgcangca cgcgtacgta agctcggaaat tcggctcgag gtgaagccac caatcatcta 60

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 ttaccaagcg cccaatgaag ggaatgctta ccggtcctgt taccattctc aactgggtcct 180
 ttgttagaaa tgaccaacct agatctgaga ccacctacca gattgctttg gctatcaagg 240
 acgaatggag gaccttgaaa aggctggcat cactgttatc caaattgatg aagctgcttt 300
 gagagagggg c 311

<210> 2855
 <211> 324
 <212> nucleic acid
 <213> Glycine max
 <400> 2855

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 cgctgatgtt atcaccattg agaactctcg ctccgatgag aagctcctgt ncagtcttcc 120
 gtgaagggtg gaagtatggg gctggaattg gccctgggtg ctatgacatc cactccccc 180
 gaataccacc aactgaagaa atcgctgaca gaatcaataa gatgcttgca gtgctcgaga 240
 agaacatctt gtgggtcaac cctgactgtg gttcaagacc cgcaagtaca ctgaagtga 300
 gccagccctt nacaaacatg gttg 324

<210> 2856
 <211> 311
 <212> nucleic acid
 <213> Glycine max
 <400> 2856

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 aagctcacta cttggactgg gctgtccatg ccttcagaat caccaatgtt ggtgtccagg 180
 ataccacca gatccacact cacatgtgct actcgaactt caacgacatc atccactcca 240
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 gtcagtcttc c 311

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 <212> nucleic acid

<213> Glycine max

<400> 2857

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taccaccaac tgaagaaatt gctgacagaa tcaacaagat gctggcagtg ctcgagaaga 180
acatcttgtg ggtgaaccct gactgtgggc tcaagaccg taagtacact gaggtgaagc 240
cagccctana aananggt 258

<210> 2858

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2858

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tggttcccggt tgcgtgaagc caccgatcat ctatggtgat gtgagccgcc caaagccaat 180
gaccgtcttc tggatcatctc tggctcagag ctttaccaag cgcccaatga agggaatgct 240
taccgggtcct gttaccattc tcaactgggc ctttggttaga aa 282

<210> 2859

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 2859

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gctgttgatc ttgttaacga gaccaagttg gatgacgaga tcaagtcattg gctagcattt 180
gctgcacaaa aaattgttga agttaacgca ttggctaagg cattgtcttg caacaaggat 240
gtggccttct tctctgctaa tgctgcagct caggcttcaa ggaagtctc tccaaga 297

<210> 2860

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2860

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 cctgttacca ttctcaactg gtcctttgtt agaaatgacc aacctagatc tgagaccacc 120
 taccagattg ctttgtctat caaggacgaa gtggaagacc ttgaaaaggc tggcatcact 180
 gttatccaaa ttgatgaagc tgctttgaga gaggggtcttc cactgaggaa atcagagcaa 240
 gctcactact tggactgggc tgtccatgcc ttcagaatca ccaatg 286

<210> 2861

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2861

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 tgaggccttc ttctctggta atgctgctgc tctggcttca aggaagtctt ctccaagagt 120
 gaccaacgag gctgtncaga aggctgctgc tgcattgagg gttcagatca tcgccgtgca 180
 acaaatgtca gtgccagact ggattctcaa caaaagaagc tcaaccttcc aatcctgcca 240
 accaccacta ttggatcctt ccctcagact gtagaactga ggaggggtacg ccgtgaattc 300
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<210> 2862

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2862

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 ctccgatgag aagctcctgt cagtcttccg tgaagggtgtg aagtatggtg ctggaattgg 120
 ccctgggtgc tatgacatcc actccccaag aataccacca actgaagaaa tcgctgacag 180
 aatcaataag atgcnnngcag tgctcgagaa gaacatcttg tgggtcaacc ctgactgtgg 240
 tctccaagac ccgcaagtac actgaagtga agccagccct cacaacatg gttgccgcag 300
 caaaactcat c 311

<210> 2863
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 2863

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 aagtatggtg ctggaattgg ccttggtgtc tatgacatcc actccccaag aataccacca 120
 actgaagaaa ttgctgacag aatcaacaag atgctggcag tgctcgagaa gaacatcttg 180
 tgggtgaacc ctgactgtgg gctcaagacc cgtaagtaca ctgaggtgaa gccagccctc 240
 acaaacatgg ttnnccgacg aaaactcatc cgcaacgaac ttgccaagtg anggta 296

<210> 2864
 <211> 305
 <212> nucleic acid
 <213> Glycine max

 <400> 2864

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 actntattgt ccctgaattg ggccctgatg tgaacttcac ctatgcttct cacaaggctg 180
 ttgatgaata caaggaggcc aaggcgcttg gagtggatac cattcccgta ctcggtggcc 240
 ctgttacata cttgttgctc tccaagcctg ccaagggagt cgagaaatcc ttttctctcc 300
 tctct 305

<210> 2865
 <211> 280
 <212> nucleic acid
 <213> Glycine max

 <400> 2865

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 actccatcat cgacatggac gctgatgtta tcaccattga gaactctcgc tccgatgaga 120
 agtcctgtc agtcttccgt gaagggtgtga agtatggtgc tggaattggc cctggtgtct 180
 atgacatcca ctccccaaga ataccaccaa ctgaagaaat cgctgacaga atcaataaga 240

tgcttgcaagt gctcgagaag aacatcttgt ggggtcaaccc 280

<210> 2866
 <211> 287
 <212> nucleic acid
 <213> Glycine max

<400> 2866

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 ctggtaaatgc tgctgctctg gttcaagga agtcttctcc aagagtgacc aacgaggctg 120
 ttcagaaggc tgctgctgca ttgaagggtt cagatcatcg ccgtgcaaca aatgtcagtg 180
 ccagactgga ttctcaacaa aagaagctca acctccaat cctgccaacc accactattg 240
 gatccttccc tcagactgta gaactgagga gggtaacgag tgaattc 287

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<210> 2867
 <211> 324
 <212> nucleic acid
 <213> Glycine max

<400> 2867

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 tgatattgat gttcttggtc atggagaacc agagagaaat gatatgggtg agtacttcgg 120
 tgaacaattg tncaggcttt gccttcaccg ttaatgggtg ggtgcaatcc tatggttccc 180
 gttgcgtgaa gccacgatca tctatggtga tgtgagccgc ccaaagccaa tgaccgtctt 240
 ctggtcattc ctggctcaga gctttaccaa gcgcccgaatg aagggaatgc ttaccggtcc 300
 tgttaccatt ctcaactggc cctt 324

<210> 2868
 <211> 273
 <212> nucleic acid
 <213> Glycine max

<400> 2868

gtcnnatgca cgcgtncgta agctcggaat tcnctcgag tggagaacca gagagaaatg 60
 atatgggtga gtacttcggt gagcaattgt caggctttgc cttcactggt aatgggtggg 120
 tgcaatccta tggttcccggt tgtgtgaagc caccaatcat ctatgggtgat gtagccgccc 180

aaagccaatg actgtcttct ggtcatctct ggctcagagc tttaccaagc gcccaatgaa 240
gggaatgctt accggtctctg ttaccattct caa 273

<210> 2869
<211> 296
<212> nucleic acid
<213> Glycine max
<400> 2869

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aaggggttcag atcatcgccg tgcaacaaat gtcagtgcc aactggatgc tcaacaaaag 120
aagctcaacc ttccaatcct tccaaccacc actattggat ccttccctca gactgtagaa 180
ctgaggaggg tacgccgtga gttcaaggct aacaagatct ccgaggaaga gtatgttaag 240
tcaattaagg aggaaattcg caaagttggt gaacttcaag aagagcttga tattga 296

<210> 2870
<211> 301
<212> nucleic acid
<213> Glycine max
<400> 2870

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agtcttctcc aagagtgacc aacgaggctg ttcagaaggc tgctgctgca ttgaagggtt 120
cagatcatcg ccgtgcaaca aatgtcagtg ccagactgga ttctcaacaa aagaagctca 180
accttccaat cctgccaacc accactattg gatccttccc tcagactgta gaactgagga 240
gggtacgccg tgaattcaag gctaacaaga tctccgagga agagtatgta aagtcaatta 300
a 301

<210> 2871
<211> 300
<212> nucleic acid
<213> Glycine max
<400> 2871

cgtcgcangc acgcgtacgt nagctcgga ttcggctcga ngcagaactt gcgcctgctt 60
tgtctgggtt gaatgttctt gttgagacct actttgctga catccttgcg gaggcataca 120

<211> 295
 <212> nucleic acid
 <213> Glycine max

 <400> 2874

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 ttgatgaagc tgctttgaga gagggctctc cactgaggan atcagancaa gctcactact 180
 tggactgggc tgtccatgcc ttcagaatca ccaatgttgg tgtccaggat accacccaga 240
 tccacactca catgtgtctac tcgaattcaa cgacatcadc cactccatca tcgag 295

<210> 2875
 <211> 303
 <212> nucleic acid
 <213> Glycine max

 <400> 2875

 nttngcacgc gtacgtaagc tcggaattcg gctcgagcctt gctgtctaca aggaagttat 60
 tgctgacctt aaggcagctg gtgcttcatt gattcagttt gatgagccta cccttgtctt 120
 ggaccttgag tctcacaagt tgcaagcatt cactgacgca tatgcagaac ttgcgcctgc 180
 tttgtctggt ttgaatgttc ttgttgagac ctactttgct gacatccctg ctgaggcata 240
 caagaccctc acatctctga atggcgctcac tgcatatgga tttgatttgg tccgtggaac 300
 caa 303

<210> 2876
 <211> 293
 <212> nucleic acid
 <213> Glycine max

 <400> 2876

 annnganac gtcgcangca cgcgtacgta agctcgggaat tcggctcgag nactcacat 60
 gtgctactcg aacttcaacg acatcatcca ctccatcadc gacatggacg ccgatgttat 120
 caccattgag aactctcgct ccgacgagaa gctnctgtca gtcttccgcy aaggtgtgaa 180
 gtatggtgct ggaattggcc ctggtgtcta tgacatccan tccccagaa taccaccaac 240
 tgangaaatt gctgacagaa tcaacaagat gctggcantg ctcgagaaga aca 293

<210> 2877
 <211> 291
 <212> nucleic acid
 <213> Glycine max

 <400> 2877

 gncgtcgcan gcacgcgtac gtaagctcgg aattcggctc gaggttagaa atgaccaacc 60
 tagatctgag accacctacc agattgcttt ggctatcaag gacgaagtgg aggaccttga 120
 aaaggctggc atcactgtta tccaaattga tgaagctgct ttgagagagg gtctgccact 180
 gaggaaatca gaacaagctc actacttgga ctgggctgtc catgccttca gaatcaccaa 240
 tgttggtgtg caggatacca ctcagatcca caccacatg tgctactcca a 291

<210> 2878
 <211> 298
 <212> nucleic acid
 <213> Glycine max

 <400> 2878

 ctnanngcgt cgcangcacg cgtacgttag ctcggaattc ggctcgaggg gaatgcttac 60
 cggctcctgtt accattctca actggtcctt tgtagaaaat gaccaacctt gatctgagac 120
 cacctaccag attgctttgt ctatcaagga cgaagtggaa gaccttgaaa aggctggcat 180
 cactgttatc caaattgatg aagctgcttt gagagagggt cttccactga ggaaatcaga 240
 gcaagctcac tacttggtgact gggctgtcca tgccctcaga atcaccaatg ttggtgtc 298

<210> 2879
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 2879

 agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gatgggtggg tgcaatccta 60
 tggttcncgt tgtgtgaagc caccaatcat ctatggtgat gtgagcngnc caaagccaat 120
 gactgtcttc tggctcatctc tggctcagag ctttaccaag cgcccaatga agggaatgct 180
 taccggtcct gttaccattc tcaactggtc ctttggttaga aatgaccaac ctagatctga 240
 gaccacctac cagattgctt tggctatcaa ggacgaagtg ggaggacctt gaaaaggctg 300

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<400>	2880
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[illegible]

<400> 2881

[illegible]

<400>	2882
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1014

gtgaacaatt gtcaggcttg ccttcaccgt taatgggtgg gtgcaatcct aggttcccgt 300
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<210> 2883
 <211> 276
 <212> nucleic acid
 <213> Glycine max
 <400> 2883

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 gagctttacc aagcgcccaa tgaagggaat gcttaccggt cctgttacca ttctcaactg 180
 gtcctttggt agaaatgacc aacctagatc tgagaccacc taccagattg ctttgtctat 240
 caaggacgaa gtggaagacc ttgaaaaggc tggcat 276

<210> 2884
 <211> 288
 <212> nucleic acid
 <213> Glycine max
 <400> 2884

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 actgtcttct ggtcatctct ggctcagagc ttaccaagc gcccaatgaa gggaatgctt 180
 accggtcctg ttaccattct caactggtcc tttgttagaa atgaccaacc tagatctgag 240
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<210> 2885
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 2885

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 cctcccttct tcacactgct gttgatcttg ttaacgagac caagttggat gacgagatca 180

agtcattggct agcatttgct gcacaaaaaa ttgttgaagt taacgcattg gctaaggcat 240
 tgtctggcaa caaggatgtg gccttcttct ctgctaattgc tgcagctcag gcttcaagga 300
 agtcctctcc aaga 314

<210> 2886
 <211> 304
 <212> nucleic acid
 <213> Glycine max
 <400> 2886

gncgcatgca cgcgtacgta agctcggaat tcggctcgag cttgatttga tcaaggggtgg 60
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 tgaccttget gcttctctca ctaccttgca gggctctgag ggcattgtgg gcaaagataa 180
 gcttgtttgtg tccacctcct cctcccttct tcacactgct gttgatctag ttaacgagac 240
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 aacg 304

<210> 2887
 <211> 275
 <212> nucleic acid
 <213> Glycine max
 <400> 2887

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 cctgttacca ttctcaactg gtcctttgtt agaaatgacc aacctagatc tgagaccacc 180
 taccagattg ctttggctat caaggacgaa gtggaggacc ttgaaaaggc tggcatcact 240
 gttatccaaa ttgatgaagc tgctttgaga gaggg 275

<210> 2888
 <211> 257
 <212> nucleic acid
 <213> Glycine max
 <400> 2888

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tgctgacatc cctgctgagg cgtacaagac cctcacatct ctgaatggcg tcaactgcata 120
 tgggtttgat ttgggtccgtg gaaccacatac tcttgatttg atcaaggggtg gatttcccag 180
 tggaaaatac ctctttgctg gagtgggtga tggaaggaac atctggggcca atgaccttgc 240
 tgcttctctc actacat 257

<210> 2889
 <211> 278
 <212> nucleic acid
 <213> Glycine max
 <400> 2889

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 tggatttgat ttgggtccgtg gaaccaacac tcttgatttg atcaaggggtg gatttcccag 180
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 tgcttctctc actaccttgc agggctcttga gggcattg 278

<210> 2890
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2890

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 gaggaaatca gagcaagctc actacttggc ctgggctgtc catgccttca gaatcaccaa 180
 tgttggtgtc caggatacca cccagatcca cactcacatg tgctactcga acttcaacga 240
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<210> 2891
 <211> 316
 <212> nucleic acid
 <213> Glycine max
 <400> 2891

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agtcaattaa ggaggaaatt cgcaaagttg ttagaacttc aagaagagct tgatattgat 120
 gttcttggtc atggagaacc agagagaaat gatatgggtg agtacttcgg tgagcaattg 180
 tnangctttg ctttactgt taatgggtgg gtgcaatcct atggttcccg ttgtgtgaag 240
 ccaccaatca tctatggtga tgtgagccgc ccaaagccaa tgactgtctt ctggtcattt 300
 ctggctcaga gcttta 316

<210> 2892
 <211> 289
 <212> nucleic acid
 <213> Glycine max
 <400> 2892

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 agaagctcaa cttccaatc cttccaacca nactattgg atccttccct cagactgtag 120
 aactgaggag ggtacgccgt gagttcaagg ctaacaagat ctccgaggaa gagtatgtta 180
 agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca agaagagctt gatattgatg 240
 ttcttggtca ggagaaccag agagaaatga tatgggtgag tacttcggt 289

<210> 2893
 <211> 320
 <212> nucleic acid
 <213> Glycine max
 <400> 2893

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 gagccgcca aagccaatga ccgtcttctg gtcattcttg gtcagagct ttaccaagcg 120
 cccaatgaag ggaatgctta ccggtcctgt taccattctc aactggctct ttgttagaaa 180
 tgaccaacct agatctgaga ccacntacca gattgctttg tctatcaagg acgaagtgga 240
 agaccttgaa aaggctggca tctactgttat ccaaattgat gaagcgcttt gagagagggt 300
 cntccactga ggaaatcaga 320

<210> 2894
 <211> 302
 <212> nucleic acid
 <213> Glycine max

<400> 2894

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cctatgggttc ccgttgctg aagccaccga tcctctatgg tgatgtgagc cgcccaaagc 120
caatgaccgt cttctgggtc tctctggctc agagntttac caagcgccca atgaaggga 180
tgcttaccgg tcctgttacc attctcaact ggtcctttgt tagaaatgac caacctagat 240
ctgagaccac ctaccagatt gctttgtcta tcaaggacga atggaagacc ttgaaaaggc 300
tg 302

<210> 2895

<211> 313

<212> nucleic acid

<213> Glycine max

<400> 2895

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tcctatgggtt ccggttgctg gaagccaccg atcatctatg gtgatgtgag ccgcccagaag 120
ccaatgaccg tcttctggct atctctggct cagagcttta ccaagcgccc aatgaaggga 180
ntgcttaccg gtctgttacc cattctcaac tggctccttg ttagaaatga ccaacctaga 240
tctgagacca cctaccagat tgctttgtct atcaaggacg aatggaagac cttgaaaagg 300
ctggcatcat gtt 313

<210> 2896

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2896

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ggagtgggtg atggaaggaa catctgggcc aatgaccttg ctgcttctct cactaccttg 120
cagggtcttg agggcattgt gggcaaagat aagcttggtg tgtccacctc ctctccctt 180
cttcacactg ctgttgatct agttaacgag accaagttgg atgatgagat caagtcatgg 240
ctagcttttg ctgccccaaa aattgttgaa gttaacgcgt ggctaaagca tgtctggcca 300
caaggatgag gg 312

<210> 2897
 <211> 291
 <212> nucleic acid
 <213> Glycine max

 <400> 2897

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 atgtgagccg cccaaagcca atgaccgtct tctgggtcatc tctgggtcag agctttacca 120
 agcgcccaat gaagggaatg cttaccggtc ctgttaccat tctcaactgg tcctttgtta 180
 gaaatgacca acctagatct gagaccacct accagattgc tttgtctatc aaggacgaag 240
 tggaagacct tgaaaaggct ggcatactg ttatccaaat tgatgaagct g 291

<210> 2898
 <211> 312
 <212> nucleic acid
 <213> Glycine max

 <400> 2898

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 aaccactcca gcnaagataa ncttgtngtg tccacctcct cctcnnttct tcacactgct 120
 gtngatnttg ttaacgagac caagttggat gacgagatca agtcattggt agcatttgct 180
 gcacaaaaaa ttgttgaagt taacgnattg gctaaggcnt tgtctggcaa caaggatgtg 240
 gccttcttnt ctgctaattgc tgcagctcag gcttcaagga agtcntctcc aagagtgacc 300
 aacgaggctg tt 312

<210> 2899
 <211> 247
 <212> nucleic acid
 <213> Glycine max

 <400> 2899

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 tgctgacatc cctgctgagg cgtacaagac cctcacatct ctgaatggcg tcaactgcata 120
 tgggtttgat ttgggtccgtg gaaccatac tottgatttg atcaagggtg gatttcccag 180
 tggaataac ctctttgctg gagtggttga tggaaggaac atctgggcca atgaccttgc 240

tgcttct 247

<210> 2900
<211> 317
<212> nucleic acid
<213> Glycine max

<400> 2900

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taccattctc aactggctct ttgttagaaa tgaccaacct agatctgaga ccacctacca 120
gattgctttg tctatcaagg acganacngg aagacctga aaaggctggc anccactggt 180
ntccaaattg atgaagctgc ttgagagag ggtcttccac tgaggaaatc agagcaagct 240
cactacttgg actgggctgt ccatgccttc agaatcacca atgttggtgt ccaggatacc 300
acccagatcc acactna 317

<210> 2901
<211> 285
<212> nucleic acid
<213> Glycine max

<400> 2901

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gatcaagtca tggctagcat ttgctgcaca aaaaattggt gaagttaacg cattggctaa 120
ggcattgtct gncaacaagg atgtggcttc ttctctgcta atgctgcagc tcaggcttca 180
aggaagtcct ctccaagagt gaccaacgag gctgttcaga aggctgctgc tgcattgaag 240
ggttcagatc atcgccgtgc aacaaatgtc agtgccagac tggat 285

<210> 2902
<211> 264
<212> nucleic acid
<213> Glycine max

<400> 2902

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gtatgtaaag tcaattaagg aggaaattcg caaagttggt gagcttcaag aagagcttga 120
tattgatgtt cttgttcatt gagaaccaga gagaaatgat atggttgagt acttcggtga 180

acaattgtca ggctttgect tcaccgttaa tgggtgggtg caatcctatg gttcccgttg 240
cgtgaagcca ccgatcatct atgg 264

<210> 2903
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 2903

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aacttcacct atgcttctca caaggctgtt gatgaatata aggaggccaa ggcgcttgga 120
gtggatacca ttcccgtact cgttggccct gttacatact tgttgctctc caagcctgcc 180
aagggagtcg agaaatcctt ttctctctc tctctccttc ccaaggttct tgctgtctac 240
aaggaagtta ttgctgacct taaggcagct ggtgcttcat ggattcaatt gatgagcct 299

<210> 2904
<211> 305
<212> nucleic acid
<213> Glycine max

<400> 2904

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ggcagctggg gcttcatggn ttcaatttga tgagcctacc cttgtcttgg accttgaatc 120
tcacaagttg caagctttca ctgacgcata tgcanaactt gcacctgctt tgtctgatct 180
gaatgttctt gttgagacct actttgctga catccctgct gaggcgtaca agaccctcac 240
atctctgaat ggcgtcactg catatgggtt tgatttggtc cgtggaaccc atactcttga 300
tttga 305

<210> 2905
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 2905

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caagcgccca atgaaggga tgcttacncc tctgtttacc attctncaac tggtcctttg 120

ttagaaatga ccaacctaga tctgagacca cctaccagat tgctttgtct atcaagntga 180
 agtggaagac cttgaaaagg ctggcatcac tgttatccaa attgatgaag ctgctttgag 240
 agaggggtctt ccaactgagga aatcagagca agctcactac ttggactggg ctgtccatg 299

<210> 2906
 <211> 286
 <212> nucleic acid
 <213> Glycine max

<400> 2906

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 tacctctttg ctggagtggg tgatggaagg aacatctggg ccaatgacct tgctgcttct 120
 ctcactacat tgcaggggtct tgagggcatt gtgggcaaag ataagcttgt tgtgtccacc 180
 tccctccct tcttcacact gctgttgatc ttgttaacga gaccaagttg gatgacgaga 240
 tcaagtcatg gctagcattt gctgcacaaa aaattgttga agttaa 286

<210> 2907
 <211> 313
 <212> nucleic acid
 <213> Glycine max

<400> 2907

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 aggcagctgg tgcttcatgg attcaatntg atgagcctac ccttatcttg gaccttgaat 120
 ctcanaagtt gcaagctttc actgacgcat atgcagaact tgcacctgct ttgtctgac 180
 tgaatgtntc ngtnagacn cactttgctg acatccctgc tgaggcgtac aagaccctca 240
 catctctgaa tggcgtcact gcatatgggt ttgatttggt ccgtggaacc catactcttg 300
 atttgatcaa ggg 313

<210> 2908
 <211> 274
 <212> nucleic acid
 <213> Glycine max

<400> 2908

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agctcaacct tccaatcctt ccaaccacca ctattggntc cttccctcag actgtagaac 120
 tgaggagggt acgccgtgaa ttcaaggcta acaagatctc cgaggaagag tatgtaaagt 180
 caattaagga ggaaattcgc aaagttgttg agcttcaaga agagcttgat attgatgttc 240
 ttgttcatgg agaaccagag agaaatgata tgggt 274

<210> 2909
 <211> 276
 <212> nucleic acid
 <213> Glycine max
 <400> 2909

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 gcccaatgaa gggaatgctt accggctctg ttaccattct caactgggtcc tttgttagaa 120
 atgaccaacc tagatctgag accacctacc agattgcttt gtctatcaag gacgaagtgg 180
 aagaccttga aaaggctggc atcactgtta tccaaattga tgaagctgct ttgagagagg 240
 gtcttccact gaggaaatca gagcaagctc atactt 276

<210> 2910
 <211> 252
 <212> nucleic acid
 <213> Glycine max
 <400> 2910

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 aatgttcttg ttgagaccta ctttgctgac atccctgctg aggcgtacaa gaccctcaca 120
 tctctgaatg gcgtcactgc atatgggttt gatttgggtcc gtggaacca tactcttgat 180
 ttgatcaagg gtggatttcc cagtggaaaa tacctctttg ctggagtggg tgatggnagg 240
 nacatctggg cc 252

<210> 2911
 <211> 286
 <212> nucleic acid
 <213> Glycine max
 <400> 2911

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gcaacaagga tgtggccttc ttctctgcta atgctgcagc ttcaggttc aaggaagtcc 120
tctccaagag tgaccaacga ggctgttcag aaggctgctg ctgcattgaa gggttcagat 180
catcgccgtg caacaaatgt cagtgccaga ctggntgctc aacaaaagaa gctcaacctt 240
ccaatccttc caaccaccac tattgntcct tcctcagacg tgtgaa 286

<210> 2912
<211> 293
<212> nucleic acid
<213> Glycine max
<400> 2912

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aacgaggctg ttcagaaggc tgctgctgca ttgaaggggt cngatcatcg ccgtgcaaca 120
aatgtcagtg ccagactgga ttctcaacan aagaagctca accttccaat cctgccaaac 180
accactattg gatccttccc tcagactgta gaactgagga gggtagcccg tgaattcnag 240
gctaacaaga tctccgagga agagtatgta nngtcaatta agnggaaaat tcg 293

<210> 2913
<211> 274
<212> nucleic acid
<213> Glycine max
<400> 2913

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ctgctgcatt gaaggggttca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatt 120
ctcaacaaaa gaagctcaac ctccaatcc tgccaaccac cactattgga tccttccttc 180
agactgtaga actgaggagg gtacgccgtg aattcaaggc taacaagatc tccgaggaag 240
agtatgtaaa gtcaattaag gaggaaattc gcaa 274

<210> 2914
<211> 283
<212> nucleic acid
<213> Glycine max
<400> 2914

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atattgnngt tcntgntcat ggaganccan agagaaatga tatggttgag tacttcgggtg 120
aacaattgtc aggcttttgct ttcaccgtta atgggtgggt gcaatcctat ggntcccgtt 180
gcgtagaagcc ancgatcatc tatggtgatg tnagccgccc aaagccaatg accgtnttct 240
ggtcattctct ggctcagagc tttaccaagc gccaatgaag gga 283

<210> 2915
<211> 534
<212> nucleic acid
<213> Glycine max
<400> 2915

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cacgcgtccg nncacnctc cgctgcagaa gacgaactgaa nggagagctt gatattgatg 120
ttcttgggca tggagaacca gagagaantg atgtggttga gtacttcgggt gagcaattgt 180
caggctttgc cttcactgtt aatgggtggg tgcaatccta tggttcccgt tgtgtgaanc 240
caccnatcat ctatggtgat gtgagcccc aaanccaatg actgtcttct ggtcattctct 300
ggctcananc tttaccaatc gcccnntgaa anggaatnct tnccgggctt gttacattnt 360
naacttgggc cttnttttna anatnancaa cctatttntt annccnctnc nttattnttt 420
tnncttttna ggatnatnng nttgnntttt tanaaanggg ttnggatnat ttntttntn 480
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<210> 2916
<211> 297
<212> nucleic acid
<213> Glycine max
<400> 2916

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gatcaacagc agtgtgaaga agggaggagg agatcaagtc atggctagct ttgctgccc 120
aaaaaattgt tgaagttaac gcattggcta aagcattgtc tggccacaag gatgaggcct 180
tcttctctng taatgctgct nctctggctt caaggaagtc ttctccaaga gtgaccaacg 240
aggctgttca gaaggctgct gctgcattga agggttcaga tcatcgccgt gcaacaa 297

<210> 2917

<211> 410
<212> nucleic acid
<213> Glycine max

<400> 2917

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tgaagaaatc gctgacagaa tcaataagat gcttgacagt ctcgagaaga acatcttctg 120
ggtcaaccct gactgtggct ttaagacccg caagtacact gaagtgaagc cagccctcac 180
aaacatgggt gccgcagcaa aactcatccg taacgaactt gccaaagtga tgggtataaga 240
aagtagaatc tacaagttca ttggttctgc ttttataata caccaaagaa aaattttcta 300
tattgggttg tttcaataac cgtgtgtgga atatttagat gttttagcat gctccgtgaa 360
caattgatcc tcctcaaacc ctctcccctt aattttccca actcccgtt 410

<210> 2918
<211> 333
<212> nucleic acid
<213> Glycine max

<400> 2918

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tcgagnnagg aggggtacgcc gtgagttcaa ggctaacaag atctccgagg aagagtatgt 120
taagtcaatt aaggaggaaa ttgcgaaagt tgttgaactt caagaagagc ttgatattga 180
tgttcttgtt catggaganc cngggngaen tgatatgggt gagtacttcg gtgagcaatt 240
gtcaggcttg ctttactgt taatgggtgg gtgcaatcca anggtcccg tgtgtnaagc 300
caccaatcca ctatggtgat gtgagccgcc caa 333

<210> 2919
<211> 313
<212> nucleic acid
<213> Glycine max

<400> 2919

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antgcttacc ggtcctgtta ccattctcaa ctggctcctt gttagaaatg accaacctag 120
atctgagacc acctaccaga ttgctttgtc tatcaaggac gaagtggaag accttgaaaa 180

ggctggcatc actgttatcc agnttgggtga agctgctttg agagaggggtc ttccactgag 240
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gtccaggata cca 313

<210> 2920
<211> 259
<212> nucleic acid
<213> Glycine max
<400> 2920

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attggctaag gcattgtctg gcaacaagga tgtggccttc ttctctgcta atgctgcagc 120
tcaggcttca aggaagtctc ctccaagagt gaccaacgag gctgttcaga aggctgctgc 180
tgcattnaag ggttcagntc ancgncgtnc aacanntcnc agccnnantg ganantcncn 240
aaaaaaggct cncncncc 259

<210> 2921
<211> 286
<212> nucleic acid
<213> Glycine max
<400> 2921

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ggccctgtta catacctgtt gctctccaag cctgccaaagg gagttgagaa atccttttct 120
ctcctctctc tccttcccaa ggttcttgct gtctacaagg aagttattgc tgaccttaag 180
gcagctggtg cttcatggat tcagtttgat gagcctaccc ttgtcttgga ccttgagtct 240
cacaagttgc aagcattcac tgacgcatac gcagaacttg cgctg 286

<210> 2922
<211> 242
<212> nucleic acid
<213> Glycine max
<400> 2922

gtgcaatcct atgggtcccg ttgcgtgaag ccaccgatca tctatgggtga tgtgagccgc 60
ccaaagccaa tgaccgtctt ctggctcatct ctggctcaga gctttaccaa gccccaatga 120

aggaatgct taccggctct gttaccattc tcaactggtc ctttgtaga aatgaccaac 180
ctagatctga gaccacctac cagattgctt tgtctatcaa ggacgaatgg aagacctga 240
aa 242

<210> 2923
<211> 270
<212> nucleic acid
<213> Glycine max

<400> 2923

gogtacgtaa gctcggaatt cggctcgagt cagatcatcg ccgtgcaaca aatgtcagtg 60
ccagactgga ttctcaacaa aagaagctca accttccaat cctgccaaacc accactattg 120
gatacttccc tcagactgta gaactgagga gggtagccg tgaattcaag gctaacaaga 180
ttctcgagga agagtatgta agtcaattaa ggaggaaatt cgcaaagttg ttgagcttca 240
agaagagctt gatattgatg ttcttgttca 270

<210> 2924
<211> 292
<212> nucleic acid
<213> Glycine max

<400> 2924

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aatactcttt tgctggagtg gttnatggaa ggaacatctn gngccaatga ccttgctgct 120
tctctacta cattgcaggg tcttgagggc attgtnggca aagataagct tgttgtgtcc 180
acctctctct ccttcttca cactgctgtt gatcttgta acgagaccaa gttggatgac 240
gagatcaagt catggctagc atttgctgca caaaaaattg tgaagttaac gc 292

<210> 2925
<211> 312
<212> nucleic acid
<213> Glycine max

<400> 2925

aanatttnnc gtcgcatgca tgcgtacgta agctcggaat tcggctcgag gagaaatgat 60
atggttctcc atgaacaaga acatcaatat caagctcttc ttgaagttct tgttcatgga 120

gaaccagaga gacnatgata tgggtgagta cttcgggtgag caattgtcag gctttgcctt 180
 cactgttaat ggggtgggtnc natectatgg ttcccgttgt gtgaagccac caatcatcta 240
 tgggtgatgtg agcgcgcccac agccaatgac tgtcttcttg tcctctcttg ctcagagctt 300
 taccaagcgc cc 312

<210> 2926
 <211> 287
 <212> nucleic acid
 <213> Glycine max
 <400> 2926

tcgcangcac gcgtacgtna gctcgggaatt cggctcgagc aagtgggttcg acaccaacta 60
 ccaactggnat tgtccctgaa ttggggccctg atgtgaactt cacctatgct tctcacaagg 120
 ctggtgatga atacaaggag gccaaggcgc ttggagtggg taccattccc gtactcggtt 180
 gccctgttac atacttggtg ctctccaagc ctgccaaggg agtcgagaaa tctttttctc 240
 tctctctctc ccttcccaag gttcttgctg tctacaagga agttatt 287

<210> 2927
 <211> 258
 <212> nucleic acid
 <213> Glycine max
 <400> 2927

aagccaccga tcctctatgg tgatgtgagc cgcccaaagc caatgaccgt cttctgggtca 60
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 attctcaact ggtctttggt agaaaatgac caacctagat ctgagacaac taccagattg 180
 ctttgtctat ccaaggacga ntnggaagac cttgaaaagg ctggcatcac tgttatccaa 240
 attgatgaag ctgctttg 258

<210> 2928
 <211> 335
 <212> nucleic acid
 <213> Glycine max
 <400> 2928

gtcgcangca cgcgtacgtn agctcgggaa ttcggctcga gctggaattg gccctgggtg 60

ctatgacatc cactcccca gaataccacc aactgaagaa atcgctgaca gaatcaataa 120
gatgcttgca gtgctcgaga agaacatctt gtgggtcaac cctgactgtg gtctcaagac 180
ccgcaagtac actgaagtga agccagccct cacaacatg gttgccgcag caaaactcat 240
ccgtaacgaa cttgccaaagt gaatggtata agaaagtaga atctacaagt tcattgggtc 300
tgcttttata atacaccaa gaaaaatttt ctata 335

<210> 2929
<211> 279
<212> nucleic acid
<213> Glycine max
<400> 2929

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CAAGGCGCTTGGAGTGGGATCCATTCCCCTACTCGTTGGCCGTGTTACATACTTGTTGCTCTCCAAGCCT
GCCAAGGGAGTCGAGAAATCCTTTCTCTCTCTCTCTCTCTTCCAAGGTCTTGCTGCTTACAAGGAAGTTATTGCTG

annntangta cgcgtacgta agctcggaat tcggctcgag caccaactac cactttattg 60
tccctgaatt gggccctgat gtgaacttca cctatgggtc tcacnaggct gttgatgaat 120
acaaggnggc caaggcgctt ggagtggata ccattcccgt actcgttggc cctgttacat 180
acttgttgct ctccaagcct gccaaaggag tcgagaaaatc cttttctctc ctctctctcc 240
ttccaaggt tcttgctgtc tacaaggaag ttattgctg 279

<210> 2930
<211> 282
<212> nucleic acid
<213> Glycine max
<400> 2930

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tcctgtcagt cttccgtgaa ggtgtgaagt atgggtgctgg aattggccct ggtgtctatg 120
acatccactc cccaagaata ccaccaactg aagaaatcgc tgacagaatc aatacgatgc 180
ttgcagtgct cgagaagaac atcttggtgg tcaaccctga ctgtggtctc aagaccgcga 240
agtacactga atgaagccag cctcacaaa catggttgcc gg 282

<210> 2931
<211> 261
<212> nucleic acid
<213> Glycine max
<400> 2931

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 ggtacgccgt gagttcaagg ctaacaagat ctccgaggaa gagtatgtta agtccaatta 120
 aggaggaaat tcgcaaagtt gttgaacttc aagaagagct tgatattgat gttcttgttc 180
 atggagaacc agagagaaat gatatggttg agtacttcgg tgagcattgt caggctttgc 240
 ctcaactgtta atgggtgggt g 261

<210> 2932
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2932

ctcccnacg tngcatgcac gcgtacgtna gctcngaat tcggctcgag gaaatccttt 60
 tctctcctct ctctccttcc caaggttctt gctgtctaca aggaagttat tgotgacctt 120
 aaggcagctg gtgcttcatg gattcaattt gatgagccta cccttgtctt ggaccttgaa 180
 tctcacaagt tgcaagcttt cactgacgca tatgcagaac ttgcacctgc tttgtctgat 240
 ctgaatgttc ttgttgagac ctactttgct gacatccctg ctgaggcgta caagaccc 298

<210> 2933
 <211> 298
 <212> nucleic acid
 <213> Glycine max
 <400> 2933

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 caagggagtc gagaaatcct tttctctcct ctctctcctt cccaaggctt ttgctgtcta 120
 caaggaagtt attgctgacc ttaaggcagc tgggtgcttca tggattcaat ttgatgagcc 180
 tacccttgtc ttggaccttg aatctcaciaa gttgcaagct ttcactgacg catatgcaga 240
 acttgcacct gctttgtctg atctgaatgt tcttgttgag actactttgc tgacatcc 298

<210> 2934
 <211> 269
 <212> nucleic acid
 <213> Glycine max
 <400> 2934

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gcaaggcgct tggagtggat 60
accattcccc tactcgttgg ccctgttaca tacttggtgc totccaagcc tgccaaggga 120
gtcgagaaat ctttttctct cctctctctc cttcccaagg ttotttctgt ctacaaggaa 180
gttattgctg accttaaggc agctggtgct tcatggattc aatttgatga gcctaccctt 240
gtcttggaac ttgaatctca caagttgcn 269

<210> 2935
<211> 261
<212> nucleic acid
<213> Glycine max
<400> 2935

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aattgctgac agaatcaaca agatgctggc agtgctcgag aagaacatct tgtgggtgaa 120
ccctgactgt gggctcaaga cccgtaagta cactgagggtg aagccagccc tcacaaacat 180
ggttgccgca gcaaaactca tccgcaacga acttgccaag tgaatggtat aagaaagtag 240
aatcttccaa gtcatttgggt t 261

<210> 2936
<211> 262
<212> nucleic acid
<213> Glycine max
<400> 2936

cgtnctnag ctcggaattc ggctcgagct tctggctcgc tctggctcag agctttacca 60
agcgcccaat gaagggaatg cttaccggtc ctgttaccat tcncaactgg tcctttgtta 120
gaaatgacca acctagatct gagaccacct accagattgc tttgtctatc aaggacgaag 180
tggaagacct tgaaaaggct ggcactcactg ttatocaaat tgatgaagct gctttgagag 240
agggtctcca ctgaggaaat ca 262

<210> 2937
<211> 280
<212> nucleic acid
<213> Glycine max
<400> 2937

acgcgtacgt aagctcggaa ttcggctcga ggtgaagtat ggtgctggaa ttggccttgg 60
 tgtctatgac atccactccc caagaatacc accaactgaa gaaattgctg acagaatcaa 120
 caagatgctg gcagtgctcg agaagaacat cttgtgggtg aaccctgact gtgggctcaa 180
 gacccgtaag tacactgagg tgaagccagc cctcacaaac atggttgccg cagcaaaaact 240
 ncatccgcaa cgaattgcc aatgatggtat aagaaataga 280

<210> 2938
 <211> 244
 <212> nucleic acid
 <213> Glycine max
 <400> 2938

angnnctca catctctgaa nggcgtcact gcatatgggn ttgatttgnt ccgtggancc 60
 catactcttg atttnatcaa ggggtggattt ccagtgga aatacctctt tgcaggagtg 120
 gttgatggaa ggaacatctg ggccaatgac cttgctgctt ctctcactac attgcagggt 180
 cttgagggat tgtgggcaaa gataagcttg ttgtgtccac ctctctctcc cttcttcaca 240
 ctgc 244

<210> 2939
 <211> 289
 <212> nucleic acid
 <213> Glycine max
 <400> 2939

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 tctggctcag agctttacca agcgcccaat gaagggaatg cttaccgggc ctgttaccat 120
 tctcaactgg tcctttgtta gaaatgacca acctagatct gagaccacct accagattgc 180
 tttggctatc aaggacgaag tggaggacct gaaaaggctg gcatcactgt tatccaaatt 240
 gatgaagctg cttgagagag ggtctgccat gaggaaatca gaacaagct 289

<210> 2940
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 2940

ncgtcgang cacgcgtacg tnagctcgga attcggctcg agggttcccg ttgcgtgaag 60
ccaccgatca tctatggtga tgtgagccgc ncaaagccaa tgaccgtctt ctggctgtct 120
ctggctcaga gctttaccaa gcgcccgaatg aagggaatgc ttaccggtcc tgttancatt 180
ctcaactggt cctttgttag aaatgaccaa cctagatctn cagaccacct accagattgc 240
tttgtctatc aaggacngt ggaagacctt gaaaaggctg gcatcactgt tatccaaatt 300
g 301

<210> 2941
<211> 295
<212> nucleic acid
<213> Glycine max

<400> 2941

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ggacgaagtg gaggaccttg aaaaggctgg catcactgtt atccaaattg atgaagctgc 120
tttgagagag ggtctgccac tgaggaaatc agaacaagct cactacttgg actgggctgt 180
ccatgccttc agaatcacca atgttggtgt gcaggatacc actcagatcc acaccacat 240
gtgtactcca antcaacgac atcatnccat ccatcanggg atngggcggc gattt 295

<210> 2942
<211> 295
<212> nucleic acid
<213> Glycine max

<400> 2942

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tctctgaatn gcgtcantgc atatgggttt gatttagtcc gtggaacca tactcttgat 120
ttgatcaagg gtggatttcc cagtggaaaa tacctctttg ctggantggt tnatggangg 180
nncatctggg ccaatgacct tgengcntct ctcachacat tncagggtct tgagggcatt 240
gtgggcnaag atannctngt tgtgtccacc tctccnccc ttentcacac tgctg 295

<210> 2943
<211> 269
<212> nucleic acid
<213> Glycine max

<400> 2943

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gctgggtgctt catggattca atttgatgag cctacccttg tcttggacct tgaatctcac 120
aagttgcaag ctttcaactga cgcataatgca gaacttgacac ctgctttgtc tgatctgaat 180
gttcttgttg agacctactt tgctgacatc cctgctgagg cgtacaagac cctcacatct 240
ctgaatggcg tcaactgcata tgggtttga 269

<210> 2944

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2944

tgcgancac gcgtacgtaa gctcggaatt cggctcgagc tgggtgtctat gacatccact 60
ccccagaat accaccaact gaagaaattg ctgacagaat caacaagatg ctggcagtgc 120
tcgagaagaa catcttgttg gtgaaccctg actgtgggct caagaccctg aagtacactg 180
aggtgaagcc agccctcaca aacatggttg ccgcagcaaa actcatccgc aacgaacttg 240
ccaagtgaat ggtataagaa agtagaatct tccaagtcac ttggtttctgc tttatattat 300
aatacaccaa ag 312

<210> 2945

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2945

gangcacgcg tacgtnagct cggaattcgg ctcgagccca gcggaaaata cctctttgct 60
gncagtgggt gatggaagga acatctgggg caatggacct ttgtggnctc tctcactac 120
cttggcaggg tcttganggg cattgtgggc aaagataagc ttgttggtgc cactcctcc 180
tcccttcttc aactgctgtg tgaccagtt aacgagacca agttggatga tgagatcaag 240
tcatggctag cttttgctgc ccaaaaaaat tgttgaagtt aacgcattgg ctaaagcatt 300
gtctggccac aaggatgagg 320

<210> 2946

<211> 299
 <212> nucleic acid
 <213> Glycine max

 <400> 2946

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 tatgggnttg atttgggtccg tggaacccat actgcttgat ttgatcaagg gtggatttcc 120
 cagtggaaaa tacctctttg ctggagtggg tgatggaagg aacatctggg ccaatgacct 180
 tgctgcttct ctactacat tgcagggctc tgagggcatt gtgggcaaag ataagcttgt 240
 tgtgtccacc tcctctctcc ttcttcacac tgctgttgat ccttggttaac gagaccaag 299

<210> 2947
 <211> 269
 <212> nucleic acid
 <213> Glycine max

 <400> 2947

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 tgacgcntat gcagaacttg gcgctgcttg ggggtggttn aatgtntctg ttgagaccga 120
 ctttntctgac atccctgctg aggcatacaa gaccctcaca tctctgaatg gcgtcactgc 180
 atatggattt gatttgggtcc gtggaaccaa cactcttgat ttgatcaagg gtggatttcc 240
 cancgaaaaa tacctctttg ctggagtgg 269

<210> 2948
 <211> 294
 <212> nucleic acid
 <213> Glycine max

 <400> 2948

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 tgtagaactg aggagggtac gccgtgagtt caaggctaac aagatctccg aggaagagta 120
 tgttaagtca attaaggagg aaattcgcaa agttgttgaa cttcaagaag agctgatatt 180
 gatgttctgt tcatggagaa ccagagagaa atgatatggg tgagtacttc ggtgagcaat 240
 tgtcaggctt tgcttcactg ttaatgggtg ggtgcatact atggttcccg ttgt 294

<210> 2949

<211> 280
 <212> nucleic acid
 <213> Glycine max
 <400> 2949
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 gtccttttgtt agaaatgacc aacctagatc tgagaccacc taccagattg ctttgtctnc 120
 caaggacgaa gtggaagacc ttgaaaagnc tggcatcaact gttatccaaa ttnatgaagc 180
 tgctttgann gaggggtcttc cactnnggaa atcagagcan ntcactactt gganccgggt 240
 gtccatntnt tcagaatcac nnntgttgggt gtccnngata 280

<210> 2950
 <211> 274
 <212> nucleic acid
 <213> Glycine max
 <400> 2950
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 acatacttgt tgctctccaa gcttgccaag ggagtcgaga aatccttttc tctcctctct 120
 ctctttccca aggttcttgc tgtctacaag gaagttattg ctgaccttaa ggcagctggt 180
 gcttcatgga ttcaatttga tgagcctacc cttgtcttgg acttgaatct cacaagttgc 240
 aagctttcat gacgcatatg cagaattgca ctgt 274

<210> 2951
 <211> 270
 <212> nucleic acid
 <213> Glycine max
 <400> 2951
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 tctcacaagt tgcaagcttt cactgangca tatgcagaac ttgcacctgc tttgtctgat 120
 ctgaatgttc ttgttgagac ctactttgct gacatccctg ctgaggcgta caagaccctc 180
 acatctctga atggcgtcac tgcataatggg tttgatttgg tccgtggaac ccatactctt 240
 gatttgatca aggggtggatt tcccagtgga 270

<210> 2952

<211> 549
 <212> nucleic acid
 <213> Glycine max
 <400> 2952

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 tcneggggtcg acccacgcgt ccggcagatc tgagaccacc taccagattg ctttgtctat 120
 caaggacgaa gtggaagacc tggaaanggc tggcatcact gttatccana ttgatgaagc 180
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 tggatcatcg canatcactg aattanaaat tttttttggt natcctnatt ttcacatatg 360
 tttgggnataa ncaantttnc gtatngacag atccannact canatgtgnc tactcggact 420
 tcaanegact ntntccaat tncattannt nancntggan tgcntgangt ntatgnnonn 480
 nttnnnannt ttntgtngna tganaagtag gttntnttn atngntatag tnnnanggtt 540
 ttnttgtn 549

<210> 2953
 <211> 317
 <212> nucleic acid
 <213> Glycine max
 <400> 2953
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 agttccaatc ctgctcaacc accacnnttg gatccttccc tacagactgt agaactgagg 120
 anggtacgcc gtgaattcna ggctaacaag atctccgagg aagagtatgt naagtcaatt 180
 aaggaggaaa ttcgcaaagt tgttgagctt caagaagagc ttgatattga tgttcttgtt 240
 catggagaac cagagagaaa tgatatgggt gagtnttcgg tgaacaattg tcaggcttgc 300
 ttcaccgtta atgggtg 317

<210> 2954
 <211> 321
 <212> nucleic acid
 <213> Glycine max
 <400> 2954

anatgcang caagcntacg taagctcgga attcggtcgc agttgngcnc tgntgtggac 60
 ttcacctatg ctttctcana aggctgtnga tgaatacnag gngggccaag ggcgcttnga 120
 gtggatacng ttccggnccct cgttggcnct gttacatagc tgttgtctctc caagcctgcc 180
 aaggagattg ngaaatccctn ttctctcctc tctctccctc ncaaggttct tgctgtctac 240
 aaggaagtta ttgntgacct taaggcagct ggtgcttcat ggattcagtt tgatgagcct 300
 acccttggtc ttggaccttg n 321

<210> 2955
 <211> 318
 <212> nucleic acid
 <213> Glycine max

<400> 2955

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 caaggctaac aagatctccg aggaagagta tgtaaagtca attaaggag gaaattcgca 120
 aagttgttga gcttcaagaa gagcttgata ttgatgttct gttcatggag aaccagagag 180
 aaatgatatg gttgagtact cgggtgaacaa ttgtcaggct tgccctaccg ttaatgggtg 240
 ggtgcaatcc tatggttccc gttgcgtgaa gccaccgatc atctatggtg atgtgagccg 300
 cccaaagcca tgaccgtc 318

<210> 2956
 <211> 260
 <212> nucleic acid
 <213> Glycine max

<400> 2956

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 ggataccgtt ccggtcctcg ttggccctgt tacatacctg ttgctctcca agcctgccaa 120
 gggagttgag aaatcctttt ctctcctctc tctccttccc aaggtcttgc tgtctacaag 180
 gaagttattg ctgaccttaa ggcagctggt gcttccatgg attcagttgg nggagctaac 240
 cctggtctgg gacctgnngt 260

<210> 2957
 <211> 247
 <212> nucleic acid

<213> Glycine max

<400> 2957

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aacaagctca ctacnttgga ctgggctgtc catgccttca gaatcnnna tnttgngng 120
cangatacna ctacgatcca caccacatg tgctactcca acttcaacga catcatccac 180
tccatcatcg acatggacgc tgatgttatc accattgaga actctcgctc cgntgagaa 240
gctcctg 247

<210> 2958

<211> 187

<212> nucleic acid

<213> Glycine max

<400> 2958

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tagatctgag accantacc agattgcttt gtctatcaag gacgangtgg aagacntga 120
aaaggctggc atcantgtna tccaaattga tgaagctgct tggagagagg gttncaccgt 180
gangaat 187

<210> 2959

<211> 250

<212> nucleic acid

<213> Glycine max

<400> 2959

aataccacca actgaagaaa ttgctgacag aatcaacaag atgctggcag tgctcgagaa 60
gaacatcttg tgggtgaacc ctgactgtgg gctcaagacc cgtaagtaca ctgaggtgaa 120
gccagccctc acaaacatgg ttccgcagc aaaactcatc cgcaacgaac ttgccaagt 180
aatgggtata ggaagtngan ttccaagtn atgggggtccg ntttaattta aaaccccccc 240
aaaaaaattt 250

<210> 2960

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2960

gtcgcangca cgcgtacgta agctcgggaat tccggtcgan ctcgagccga atcggctcga 60
gataccacca actgaagaaa ttgccgacag aatcaacaag angtctggcag tgctcgagaa 120
gaacatcttg tgggtgaacc ctgactgttg gctcaagacc cgtaagtaca ctgaggtgaa 180
gccagccctc acaaacatgg ttgccgcagc aaaactcatc cgcaacgaac ttgccaaagt 240
aatggtataa gaaagtagaa tcttccaagt catttggttc tgctttatat tat 293

<210> 2961

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2961

cnagnattca ctgacgcata tgnagaactt gcgccacgct ttgtctgggt tgacatgttc 60
ttnttgagac ctactttggc tgacatccct gctgaggcat acaagaccct cacatctctg 120
aatggcgtca ctgcatatgg atttgatttg gtccgtggaa ccaanactct tgatttgatc 180
aaggggtggat ttcccanggg aaaatacttt tttgggggan tgntgatgga aggancattg 240
ggccaatgac tttgctgttt t 261

<210> 2962

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2962

ngtcgcatgc acgcgtacgt aagctcgga ttcggnctga gggagtggat accgttccgg 60
tcctcgntgg cctctnaca tacctgttgc tntccaagc ctgccaagg agttgagaaa 120
tcctttctct cctctntn tnctcgnaagg ntnttgngt ctanaaggaa gtnattgntg 180
accttaaggc agctgggtgc tcatggattc agtttgatga gcctaccctt gtcttgacc 240
ttngtctca caagttgcaa gcattcagtg ccgcana 277

<210> 2963

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2963

agactgtaga actgaggagg gtacgccgtg agttcaaggc taacaagatc ctccgaggaa 60
gagtatgtta agtccaatta aggaggaaat tcgcaaagtt gttgaacttc aagaagagct 120
tgatattgat gttcttgttc atggagaacc agagagaaat gatatggttg agtacttcgg 180
tgagcaattn tnaggctttg cttcactgt taatggntgg gtgcantcca tggttcccgt 240
tgtgtgaagc aacaatnnac caaggnnatt aaccgcacca aagccattga ott 293

<210> 2964

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2964

accaactgaa gaaattgctg acagaatcaa caagatgctg gcagtgctcg agaagaacat 60
cttgtgngtg aaccctgact gtgggtcaa gaccgcgaag tacactgagg tnaagccagc 120
cctcacaaac atggttgccg cagcaaaact catccgcaac gaacttgcca agtgaatggc 180
ataagaaagt agaatcnnac caagtcattt gggtctgctt tatatta 227

<210> 2965

<211> 290

<212> nucleic acid

<213> Glycine max

<400> 2965

ntcgangca cgcgtacgta agctcggaat tcggctcgag gtgaccaacg aggctgttca 60
gaaggntgct gctgcattga agggtnacga tcatcgccgt gcaacaaatg tcagtgccag 120
actggattct caacaaaaga agctcaacct tccaatcctg nccaaccacc actatnggat 180
ccttcctca gactgtagaa ctgaggaggg naggcngaa ttcaaggcta acaagatctc 240
cgaggaagag tatgtaaaagt caattaagga ggaaattcgc aaagttgttg 290

<210> 2966

<211> 256

<212> nucleic acid

<213> Glycine max

<400> 2966

aagacctcac atctctgant gtnctcant gcatatgggt ttgatttggt ccgtggaagc 60
 atactcttga tttgatcaag ggnngatttc ccagtggaga atacctcttt gctggagtgg 120
 ttgatggaan gaacatcngg gccaatgacc ttgtgtcttc tcncaactaca tgcaggggtct 180
 tgagggcatt gtgggcaaag ataagcttgt tgtgnccacc tcnnoctccc ttctcacact 240
 gctgtngatc ntgtna 256

<210> 2967
 <211> 330
 <212> nucleic acid
 <213> Glycine max
 <400> 2967

gtcgcattgca cgcgtacgta agctcggaat tccgctcgan notcgaaccg ctcgagcggc 60
 tcgancgggc tcgagatcca ctccccactg aataccacca actnctagan attncctgac 120
 agaatcaaca agatgcacgc agtgctcgag aagaacatct tgtgggtgaa ccctgactgt 180
 gggctcaaga cccgtaagta cactgagggtg aagccagccc tcacaaacat ggttgccgca 240
 gcaaaaactca tccgcaacga acttgccaag tgaatgggat aagaaagtag aatcttccaa 300
 gtcatttggt tctgtttata ttataataca 330

<210> 2968
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 2968

tcgcangcac gcgtacgtaa gctcgggttc gacaccanct acnacatnaa ttgtccctga 60
 attgggccct gatgtgaact tcacctatgn cnotcacaag gctgttgatg aatacaagga 120
 ggccaaggcg cttggagtgg ataccgttcc ggtcctcgnt ggccctgtta catacctgtt 180
 gctctccaag cctgccaagg gagtnagaaa gccntttcgc tccnctctct ccggcccaag 240
 gacttgctgt cnacaaggaa gnnattgcng accngaange agcnggngca tcanggatca 300
 gttnga 306

<210> 2969
 <211> 215
 <212> nucleic acid

<213> Glycine max

<400> 2969

tnagctcgga attcggtcgc agcanggcgt tggagtggat accgttcggg tccctcgttg 60
ccctgttaca tacctgttgc tccctccaaag cctgccaagg gagttgagaa atccctttct 120
ctcctctctc tccctcccaa ggttcttgcgt gtctacaagg aagttattgc tgaccttaag 180
gcagctggtg cttcatggat tcagtttgat gagcc 215

<210> 2970

<211> 172

<212> nucleic acid

<213> Glycine max

<400> 2970

ngtcgcacgc acgcgtacgt aagctcggaa ttcggctcga gcgcacatgc agaacttgcg 60
cctgctttgt ctgggttgaa tgttcttgtt gagacctact ttgctgacat ccctgctgag 120
gcatacaaga ccctcacatc tctgaatggc gtcactgcat atggatttga tt 172

<210> 2971

<211> 170

<212> nucleic acid

<213> Glycine max

<400> 2971

gtcgcangca cgcgtacgta agctcggaa ttcggctcga ngcttctcac aaggctgttg 60
atgaatacaa ggaggccaag gcgcttggag tggataccgt tccggctctc gttggccctg 120
ttacatacct gttgctctcc aagcctgcca agggagttga gaaatccttt 170

<210> 2972

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2972

gtcgcangcn cgcgtacgtn agctcgtcaa ttcggctcga gnacttgtn gctctccaag 60
cctgnccaag ggagtcgaga aatccctttt ctctctctc tctccttccc aaggttcttg 120
ctgtctacaa ggaagttatt gctgacctta aggcagntgg tgcttcatgg attcaatttg 180

atgagcctac cctgtctgga ccttgaatct cacaagttgc aagctttcac tgacgcatat 240
gcagaacttg gcacctgctt tgtctgatct gaatgttctt gtngagacct atcntgctga 300
catccctggt gngnggtana a 321

<210> 2973
<211> 236
<212> nucleic acid
<213> Glycine max

<400> 2973

tacaaggagg ccaaggcgct tggacgtgga taccgttccg gtccctgcgtt ggccctgtta 60
aatacctggt gctctccang cctgncaang gagttgagaa atccttnnct ctccctctctc 120
tccttcccaa ggttcttgct gtctacaagg aagttattgc tgaccttaag gcagtgggtgc 180
ttcatggatt cannttnatg agtctacnct gtnttggact tgagtctcac aagttg 236

<210> 2974
<211> 231
<212> nucleic acid
<213> Glycine max

<400> 2974

actcnanncn cncntncgtn agcncgggnt tcggctcnag nttggntcnt tcnctcagac 60
tgtanaacng ngnagggtac gccgtnaatt caaggctaac aanatctgcn gnggangagt 120
atntaaagtc aattanggag gaaattcgca aagttgttga gcttcaagaa gagcttgata 180
ttgatgttct tgttcatgga gaaccagcga nanatgntat ggttnagtcc c 231

<210> 2975
<211> 313
<212> nucleic acid
<213> Glycine max

<400> 2975

tnntngcacg cgtacgtaag ctcggaattc ggctcgagat gttgtaatat ttattctgct 60
gttactcatg gcttcttttc tttcctctca ggctgctgca ttgaagggtt cagatcatcg 120
ccgtgcaaca aatgtcagtg ccagactgga ttctcaacaa aagaagctca accttccaat 180
cctgnccaac caccactatt ggatccttcc ctcagactgt agaactgagg agggtagcgc 240

ggaattcaag gctaacaaga tctccgagga agagtatgta aagtcaatta aggaggnnat 300
tcgcaaagtt gtt 313

<210> 2976
<211> 184
<212> nucleic acid
<213> Glycine max

<400> 2976

atncagcgta cgtaagtcgg aattcggctc gagcagagtt taccaagcgc ccaatgaagg 60
gaatgcttac cggctctggt ancattcttc aactggctct ttgtagaaa tgaccaacct 120
agatctgaga nccantacca gattgctttn ggctatcaaa gacgaantng agggnncttg 180
aaaa 184

<210> 2977
<211> 314
<212> nucleic acid
<213> Glycine max

<400> 2977

nngtngcang cacgcgtacg taagctcgga attcggctcg aggggtgaacc ctgactgtgg 60
gctcaagacc cgtaagtaca ctgaggtgaa gccagccctc acaaacatgg ttgccgcagc 120
aaaactcatc cgcaacgaac ttgccaagtg aatgggtataa ganagtagaa tcttccaagt 180
catttggttc tgctttatat tataatacac caaagaaaaa ttttctctat attgggttgt 240
ttcaataact gtgtgtggaa tatttaggtg tcttagcatg ctctgtgagc aattgattct 300
tcctcaaccc ctcc 314

<210> 2978
<211> 153
<212> nucleic acid
<213> Glycine max

<400> 2978

ncgcatgcac gcgtacgtaa gctcggaatt cggctcgagg tgaccaacga ggctgttcag 60
aaggctgctg ctgcattgaa gggttcagat catcgccgtg caacaaatgt cagtgccaga 120
ctggatttct caacaaaaga agctcaacct tcc 153

<210> 2979
 <211> 280
 <212> nucleic acid
 <213> Glycine max

 <400> 2979

 gtcgctgcnc gcgtaagtcg gctcgnngtt cggntcgnng ttattgctga ccttnaggca 60
 gctgggtgctt catggattca gtttgangag cctnnncttn tnttggaccc ngngnanaa 120
 ngnnnganga nnncgngag gaanatggga attntgcgcg tacttngnnn ngmntgaang 180
 annntnatna naccctctnt nntggaatac cnnatngngn aaccnngaac cctnnecatct 240
 ctggaatggc gnnatgcgta tggattgatn agtcngtgga 280

<210> 2980
 <211> 102
 <212> nucleic acid
 <213> Glycine max

 <400> 2980

 cccccagtgg anaatacctc nttgctggag tggttgatgg aaggaacatc tgggccaatg 60
 accttgctgc ttctctccac tacattgcag ggtcttgagg gc 102

<210> 2981
 <211> 288
 <212> nucleic acid
 <213> Glycine max

 <400> 2981

 gtcgangtgc acgcgtacgt aagctcggaa ttcggctoga gccacagtca gggttcaaga 60
 cccgtaagta cactgaggtg aagccagccc tcacaaacat ggttgccgca gcaaaactca 120
 tccgcaacga acttgccaag tgaatggtat aagaaagtag aatcttccaa gtcatttggt 180
 tctgctttat attataatac accaaagaaa aattttcccc atattgggtg nttnataac 240
 tgngngtgga atatttangt gncttagcat gctctgtgag caattgat 288

<210> 2982
 <211> 260
 <212> nucleic acid
 <213> Glycine max

<400> 2982

agctcggaat tcggctcgag cgtaagtaca ctgaggtgaa gccagccctc acaaacatgg 60
ttgccgcagc aaaactcctc cgcaacgaac ttgccaaagt aatggtatga gaaagtagaa 120
tcttccaagt catttggttc tgctttatat tataatacac aaagaaaaat tttctctata 180
ttgggttggt tcaataactg tgtgtggaat atttaggtgt cttagcatgc tctgtgagca 240
attgattctt cctcaacccc 260

<210> 2983

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 2983

gtcgngcac gcgtacgtaa gctcggaatt cngctcgagc totgcactct ctctttctca 60
tcctcttctc ttctgttctc tgttcgtgcc acttcttctc cgagcaatgg gcatctcata 120
ttgttggtta tccacgcatg ggaccaaga gagaacttaa gtttgctttg gaatcttttt 180
gggatggaaa gagtagtgct gatgatctgc agaaggttgc tgetgacctt aggnacagcca 240
tctggaagca gatggctgat gctggaataa agtatattcc tagcaacact ttctcatact 300
atgatcaagt actggacaca acn 323

<210> 2984

<211> 335

<212> nucleic acid

<213> Glycine max

<400> 2984

ngtcgcangc acgcntacgt nagctcggaa ttccgctcgn nctcgtctct cactctctct 60
ttctcatcct attctcttcg cttctctggt cgtgccactt cttctnccag caatggcatc 120
tcatattggt gggtatccac gcatgggacc caagagagaa ctttaagtttg ctttggaatc 180
tttttgggat ggaaagagta gtgctgagga gctgcagaag gttgctgcag accttaggtc 240
agccatctgg aagcagatgg ctgatgctgg aataaagnat attcctagca acaccttctc 300
actttacgat caagtatgga cacaacagcc atgct 335

<210> 2985

<211> 297
 <212> nucleic acid
 <213> Glycine max

<400> 2985

ngtngcatgc acgcgtacgt aagctcggaa ttcggctcga ggtttacgat ctcaactctgc 60
 actctctcct tctcatcctc ttctcttcgc ttctctgttc gtgccacttc ttctcgagca 120
 atggcatctc atattgttgg ttatccacgc atgggaccca anagagaact taagtittgct 180
 ttggaatctt tttgggatgg aaagagtagt gctgatgac tgcagaaggt tgctgctgac 240
 cttagggtcag ccatctggaa gcagatggct gatgctggaa taaagtatat tcctagc 297

<210> 2986
 <211> 327
 <212> nucleic acid
 <213> Glycine max

<400> 2986

nnttancaag cgtangtaag ctcggaattc ngctcgagct cactctgcac tctctcctct 60
 ctcatectcg tgtctncgn ncnctcggnc ngcnattccg tcccagacna tggcatctca 120
 tattgtnggt tatccacgca tggccnccca ngagaganct taagttnngct ttggaatctt 180
 tttgggatgg aaagagtagt gctgatgac tgcngaaggt tgctgctgac cttagggtcag 240
 ccatctggaa gcagatggct gatgctggaa taaagtatat tcctagcaac actttctcat 300
 actatgatca agtactggac acaacag 327

<210> 2987
 <211> 315
 <212> nucleic acid
 <213> Glycine max

<400> 2987

cnatcgcang cangcgtacg taagctcgga attcggctcg aggtgaagtg antagtataa 60
 atntacaaan tcnctncaan tcattgtgtg gcgccgttna cattctcgnt ctgcactctc 120
 tctttctcat cctnntctct tcgcntctct gtncgtgccca cntcttctcg agcaatggca 180
 tctcatattg ttggttatch acgcatggna cccaagagag aacgtaagtt ngcttnggan 240
 tctntttggg atggaaagag nagtgctgag gagctgcaga aggttgctgc agaccttagg 300

tgagccatct tgaan 315

<210> 2988
<211> 306
<212> nucleic acid
<213> Glycine max

<400> 2988

agtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggattattaa tcgtatcagt 60

gaaagaagta agaagagaga gaagtgaagt tagtagtata aatgtacaaa ctctcctcta 120

ttcagtggtg ggcgccgttt acgatctcac tctgcaactct ctctcttctca tctctctctc 180

ttcgcttctc tgttcgtgcc acttcttctc gagcaatggc atctcatatt gttgggttatc 240

cacgcatggg acccaagaga gaacttaagt ttgctttgga atcttttttg gatggaaaga 300

gtagtg 306

<210> 2989
<211> 264
<212> nucleic acid
<213> Glycine max

<400> 2989

gtcgcangca cgcntacgtn agctcggaa ttcggctcga ctctgcactc tctctttcnc 60

atcctattct cttcgcttct ctgtncogtg ccacttcttc tcgagcaatg gcatctcata 120

ttgttggtta tccacgcatg ggaccaana gagaacttaa gtttgctttg gaatcttttg 180

gatggaaaga gtagtgctga ggagctgcag aaggttgctg cagaccttag gtcagccatc 240

tggaagcaga tggctgatgc tggc 264

<210> 2990
<211> 316
<212> nucleic acid
<213> Glycine max

<400> 2990

tacgcgtacg taagctcga attcggctcg agcttctttc cctaattgac aagatccttc 60

ctgtctacag ggaggntggt gctgaattga aggcagctgg tgctacttgg atccagtttg 120

atgaacctac ccttgatga gatctcaana cccaccagtt acaagcattt acacatgnct 180

atgcagagct agagtcaagt ttatctggtt ttaatgttct gattgagana tactttgctg 240
atgtccctgc tgaagcatac aaaacactca cctctttgaa ggctgttact gcatatgggt 300
ttgatattgt tcgtgg 316

<210> 2991
<211> 321
<212> nucleic acid
<213> Glycine max
<400> 2991

caanatacat gcacgcntac gtnagctcgg aattcggctc gagggcagct ggtgctactt 60
ggatccagtt tgatgaacct acccttgtga aggatctcaa caccaccag ttacaagcat 120
ttacacatgc ctatgcagag ctagagtcaa gtttatctgg ttttaatgtt ctgattgaga 180
catactttgc tgatgtccct gctgaagcat acaaaacact cacctctttg aaggctgtta 240
ctgcatatgg gtttgatatt gttcgtggaa caaagaccct ggatttggtc naggcagggt 300
ttccncggg gaaatntttt t 321

<210> 2992
<211> 331
<212> nucleic acid
<213> Glycine max
<400> 2992

ttgcangcac gcgtagtaag ctcggaattc ggctcgagnn tctttcccta attgacaaga 60
tccttcctgg tctacagga ggtngttgct gaattgaagg cagctggtgc tatttggtac 120
tagtntgatg aacctacnt tgtgaaggat ctcaacaccc accagttaca agcatttaca 180
catgcctatg caganctaga gtcaagttta tctggtttta atgttctgat tgagacatac 240
tttgctgatg tcctgtctga agcatacaaa aactcacct ctttgaaggc tgttactgca 300
tatgggtttg atattgttcg tggaacaaag a 331

<210> 2993
<211> 284
<212> nucleic acid
<213> Glycine max
<400> 2993

gatcgacatg cttgggctgg agaatacccc ggcgtgaagc gcataccat caagcccaaa 60
 actgacagat ggggtctccct gagaccaaca cggatatcatg tcttggctga gggtcgattg 120
 atgaacttgg atgcgcaatg gaaaccccag tttggagtct gttcctcaac naccagtcac 180
 tgtcacttag tgtgaagggg natacgnagt acgagagagt tagttgccca gactgagaga 240
 gtgtgcntta ctggcaattg gactagtgcc actacagccg gtgt 284

<210> 2994
 <211> 297
 <212> nucleic acid
 <213> Glycine max

<400> 2994

cangcaecgg taogtaagct cggnatctcg ctcgaggggc cccgnacggc ggccccgacc 60
 tnategtcga cgacggagge gacgccaccc tctcatcca cgagggcgte aaggccgagg 120
 agctctatga gaagaccggg gaactccccg accctaactc cactnanaat nnaannntc 180
 cagatcgtgc ttaccnncan cagnganngg ttgaanaccg atcncaccan gtaccgnaag 240
 atgaaggncg gtctcgttgg ggtttctgag aaaccaacac tgggtgtaag agnctat 297

<210> 2995
 <211> 318
 <212> nucleic acid
 <213> Glycine max

<400> 2995

ngtcgcatgc acgcntacgt aagctcggaa ttcggctcga gtgagaagac cggggaactc 60
 cccgacccta actccactga caacgccgag ttccagatcg tgcttaccat catcagagat 120
 gggttgaaga ccgatccac caggtaccgc aagatgaagg agcgtctcgt tggggtttct 180
 gaggaanaac aaccactggt gttaaagang gctaaatccg gatgncaggn gaatggggat 240
 tctantcntt ccctgnngat aaatgtcaat gagannctnt tcnnaaggn ccangtttgn 300
 aaanttttna cgggnccg 318

<210> 2996
 <211> 288
 <212> nucleic acid
 <213> Glycine max

<400> 2996

cgtcgcangc acgcgtacgt aagctcggaa ttgggtcga gcccgacccc anctccaccg 60
ncaacgccga gtttcanatc gtgcttacna tcatcagaga tgggttgang accgatccca 120
ccaggtaccg caagatgncg gngcgtctcg ttgggggttnc tgaggnnacc accactggng 180
tnangaggct ctatnagatg naggcgaatg ggncctctct ctccctgcn attnntgtna 240
atgacnngtc nccangagca agtttgacaa cntgtatngg tgccgtca 288

<210> 2997

<211> 334

<212> nucleic acid

<213> Glycine max

<400> 2997

cgtcgcngca cgogtacgtn agctcggaa ttgggtcga tcaacatcaa gcctcaaaca 60
gacagatggg ttttccccga taccaagagg gggatcatcg tgttggcaga gggtcgtttg 120
atgaacttgg ggtgtgccac gggacacccc agctttgtga tgtcgtgctc cttcaccaac 180
caggtcattg ctcagcttga attgtggaaa gagaagggtt ctgggaagta tgagaagaag 240
gataatnnat nccaagcac ctnacgagaa agnngcntna nanctaccat tgccagcatt 300
gagcnntgcg naccaagcgt tccaaagacc aagc 334

<210> 2998

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2998

gcacgcgtac gtaagctcgg aattcggctc gagcggagtt ccagatcgtg ctgagcatca 60
tcagggatgg cttgaagacc gatcccaaga ggtaccacaa gatgaaggac agaatcgtcg 120
gtgtctccga agaaaccacc accggtgtca agaggctcta ccagatgcag gccaatggct 180
ccctcttgtt ccctgccatc aacgtcaatg actcggtcac caagagcaag tntgataact 240
tgtatggatg ccgtcactcg cttcccgatg gactgat 277

<210> 2999

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2999

tcgcangcac gcgtacgtaa gctcgnnaat tcggtctgag ctcagccctt caagggggcc 60
cgcatcaccg gctcccttca catgaccatc cagaccgtg tctcatcga gaccnncanc 120
gttctngggc ncgtgggttc ntngtgetcc tnnaanntnt tcnccattna gnnnncanc 180
gengtcgna tcgncgtnt cagcgtccgc tgtctacgcc tgtaatggtg ataccotcca 240
tgagtactnt tgggtgcaeng atnggccctc gatgggnccc cggcggcngn ccc 293

<210> 3000

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 3000

gcgaangcat gcacgcgtac gtaagctcgg aattcggctc gaggccagct tggagctagg 60
ctcaccaagc tttccaaaga ccaagctgat tacatcagtg tgctgttga gggtcctaac 120
aagcctctc actacaggtc ctgatccatc ctattnnngg agaataaacc taaactattn 180
tatcaattcc cgaggcntca ttgttacact ttcttttttg gattttttcc attacaattt 240
acntttgtgg tagcatcgga gcttcttttt tcttttttag tannatca 288

<210> 3001

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3001

gcangcacgc gtacgtaagc tcggaattcg gctcgaggtc tccgaggccg acatcttcgt 60
caccaccaca gggaacaagg acattatcat gcttgaccac atgaagaaga tgaagaacaa 120
tgccatcgtg tgcaacattg gccacttcga caacgaaatc gacatgctgg ggcttgagac 180
ctgccctggg gtgaagcgca tcaacatcaa gcctcanacn gacagatggg ttttccccga 240
taccaagagg gggatcatcg tgttggcaga gggctgtttg atgaan 286

<210> 3002

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 3002

acgncgcang gacncgtaag ntcagggtctc nagctcgnng cnaggatccc aagaggtacc 60
acaacgnnga aggacagaat cgtcgggtgtc tccgaagana ccaccaccgg tgtcaagagg 120
ctctaccaga tgcnggcnaa tggctccctc ttgttccttg ccatcaacgt caatgactcg 180
gtcaccaana gnangtttga taacttgtnt ggatnccgtc actcgtttcc cgatggactg 240
atgagagcca ctgatntgat gattgccgga aaggnagctg ttgtgtgtgg ctac 294

<210> 3003

<211> 256

<212> nucleic acid

<213> Glycine max

<400> 3003

gtcacacgag gtcccgatag antgatgaan cgccactgat gtgatgatng ccggaaaggt 60
agcagttgng gnnggcaccg agangnggca aggntagnnc ancgcnana ngnaaaccng 120
ggcncgtgtc atngtcaneg agatngatcc catntntncc cttnaggcnc taatgncggg 180
tcttcaggtt ctcacctgg aagatgtngt ctccgaggcc gacatntcg tcaccancac 240
agggaacaag gacatt 256

<210> 3004

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 3004

gtngcannca cgcgtacgta agctcggat tccgctcgaa ngggccccgg cggcggcccc 60
gacctcatcg tcgacgacgg aggcgacgcc anctcctca tccacgaggg cgtcaaggcc 120
gaggagctct atgagnagac cggggaantc cncgnnccta antccactna caacgccgag 180
ttccagattg tgcttaccat catcagngat gggttnnaana angatccan naggtaccgn 240
aanntgaagg ancgctctcg tggggtttct gangaaccga cgatgggtgtt aagatgtana 300
nc 302

<210> 3005

<211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 3005

 aaanontcaa tgcangcac gcgtacgtna gctcggaatt cggctcgagc agagatgggt 60
 tgaagaccga toccaccagg taccgcaaga tgaaggagcg tctcgttggg gttnctgagg 120
 aaaccaccac tgggtgtaag aggctatata agatgcagcg attgggatcc tatntttccn 180
 ngnaattaat gtcaatgact ctgttaccaa gancaatttg acaacttgta ccgggtgccg 240
 tcantctctc cctgatggnc tantnaggcc tactnntgtg ntgttgntcg gaaagttgnn 300
 ngtgttgcnt gat 313

<210> 3006
 <211> 306
 <212> nucleic acid
 <213> Glycine max

 <400> 3006

 agtcgcatgc acncgtacgt aagctcggaa ttccggctcga gggncgcgat caacggctcc 60
 ctccacatga cnatccagag gaggcgttnt cattgagann ntcanngncc ttggngncga 120
 ggtncgntgg tgctcntgca anatcttctc caccaggan cagcnnacg cngctattgc 180
 ccgcganagt gcngccgtct tcgntcggaa gggtagagacc ctccaggagt actggtggtg 240
 caccgagcgc gcnetcgant ggggccccgg tggtaggaccg anctcatcgt cgacgaggtg 300
 gtgang 306

<210> 3007
 <211> 70
 <212> nucleic acid
 <213> Glycine max

 <400> 3007

 attnttntgc ttgaccaca innagaanat gaagnannan tgccttggtg ttgcaacatt 60
 tggccacttt 70

<210> 3008
 <211> 536
 <212> nucleic acid

<213> Glycine max

<400> 3008

gggggngnnn nnnnttcann ggttnctgc cgtaacggtc cgaaatcccg ggtcgnccca 60
cgcgctccgcc cacgcgtccg nccgntgcga naagannaca gaagggggcc caggctgatt 120
acatcantgt gcctgttgag ggtccatnca agcctgctca ntacnngtac taagtaattg 180
agattatcaa cggaacagtg aggganagac ntantcgggtt ttatgaatcg ggntgattgt 240
ttaagtnttc cttttttttg aggttttgtt gttanacttt tcagatttga gggtagcctn 300
agtttanctt tngggcngcn naagnagnag tcaggtnttn aaaaaaggng gcngngntgg 360
nngatcnaa nttacgtacg cttgcntnca acgtcatnnc tcttcgaaag tggcaccnat 420
tttcaattca ggggcgggnc gttttaannn cnnctttnnc ggggaaaacc ttggggntan 480
ccanggttac ccccttgnen tnanncccn ttttcccnna nttgggttaa aaaaaa 536

<210> 3009

<211> 330

<212> nucleic acid

<213> Glycine max

<400> 3009

gtccnatncc gctgngngna gacnggggt tgggtnacnt ggnntcttc ntggttgncc 60
ncttgagggn cttgatganc aatgccattg tttgctncat tggtcacttt gggcctgaga 120
tngacntgct tggntcggag nactattccg gtgtgntnng catcaccatc atgnccctt 180
tctgacagat gggctctccc tgataccngc nccggtatcn ttgtcttggc tgagggtcnn 240
ttgntnatct tgggatncnc cattngncac ttcctttttg tgatgtcctg ctccttcngc 300
antngntcn ntgctcngnt tnggttgg 330

<210> 3010

<211> 473

<212> nucleic acid

<213> Glycine max

<400> 3010

gtttgattcc gagccannca atgcatcgna nncangcgta cntaaactcg gaattcggcn 60
cnagcaatga ctctgtcacc aagagcaant ttgacanctt gtatgggtgc cgtcactctc 120

tccctgatgg nctcatgagg gctaccgatg ttatgattgc tggaaaggtg gctgttgtgg 180
 ctggatatgg tgatgttggc aagggttgtg ctgctgcaat gaagcaggct ggtgctcgtg 240
 tcatcgtgac cganattgat cccatctgtg cccttcaggc tctcatggaa ggccttcagg 300
 ttctgacctt ggaggatgtt gtttctgagg ctgatatctt tgtcaccacc accggttaaca 360
 aggacatcat catggttgac cacatgagga aaatgaacaa caatgccatt gtttgcaaca 420
 ttgggtcaent tgacaatgag atcgacatgc ttgggctgga gaactaaccg ggg 473

<210> 3011
 <211> 500
 <212> nucleic acid
 <213> Glycine max

<400> 3011

gaggnntgnt ttgnanchnan anactttcgc ctgccgtacc ggtccggaat tcccgggtcg 60
 acccacgcgt ccgcggacgc gtgggcggac gcgtgggctg cgagagacga cagaaggggg 120
 gactctactc ttccctgcta ttaatgtcaa tgactctgtt accaagagca agtttgacaa 180
 cttgtacggg tgctgtcact ctctccctga tggctgatg agggctactg atgtgatgat 240
 tgctggaaaag gtggctgttg tggctggata tggngatgtt ggcaagggtt gtgctgctgc 300
 attgaagcag gctgggtgctc gtgtcatcgt gactgagatt gaccccatth gtgcccttca 360
 ggctctcatg gaaggccttc aggttctgac cttggaggat gttgtttctg aggctgatat 420
 ctttgtcacc accacgggta acaaggacat catcatggnt gaccacatga agaaaatgaa 480
 gaacaatgcc attgtttgca 500

<210> 3012
 <211> 383
 <212> nucleic acid
 <213> Glycine max

<400> 3012

ccattgtttg caacattggt cactttgaca atgagatcga catgcttggg ctggagaact 60
 accooggcgt gaagcgcac accatcaagc cccaaactga cagatgggtc ttccctgaga 120
 ccaacaccgg tatcattgtc ttggctgagg gtcgattgat gaacttggga tgcgccactg 180
 gacaccccag ttttgtgatg tcctgctcct tcaccaacca ggtcattgct cagcttgagt 240

tgtggaagga gaagagtacc ggcaagtacg agaagaaggt ttacgttttg cccaagcacc 300
 ttgatgagaa ggtggctgca cttcacctgg gcaaacttgg agctaagctg acccagctta 360
 gcaagtccca ggctgattac atc 383

<210> 3013
 <211> 528
 <212> nucleic acid
 <213> Glycine max

<400> 3013

gnnnggaggt ttganngggg gngggnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnagagagan agagagagat ctatctatct 120
 atcaagatgg ngttgttggg tgagaaaaca agcagtggaa gggagtacaa ggtgaaggac 180
 atgacgcaag ccgatttcgg aagattggaa atcgagctgg cggagggtga aatgcccggc 240
 ctcatgtect cccgcaccga gttcggcccc tctcaacct tcaagggcgc taggatcacc 300
 ggctccctcc acatgaccat ccaaaccgcc gtccctcatg agaccctcac cgccctcggc 360
 gccgaggtcc gctggtgctc ctgcaacatc ttctccaccc aggaccatgc cgccgccgcc 420
 atngcccgcg acagcgccctt cgtcttcgcc tgggaaggggt gagaccctnc aggaatactg 480
 gtggtgcacc gagcgcgccc tcgactgggg ccccggcggg gggcccca 528

<210> 3014
 <211> 520
 <212> nucleic acid
 <213> Glycine max

<400> 3014

agngtttttg tanntggggg gggggggggg aagnganata tnttagctat agatnlnaca 60
 tgtacanngt acgtaagctc ggaattcggc tcgagggccg acttcggccg cctcgagatc 120
 gagctggccc gaggttgaga tgccgggcct catggcctgg cggaccgagt tcggcccatc 180
 tcacccttca agggggcccg catcaccggc tcccttcaca tgaccatnca gaccgctgtc 240
 ctcatogaga cctcaccgn tctcggcgcc gaggttcgct ggtgctcctg caacatcttc 300
 tcaactcagga ccacgccgcc gncgccatcg cccgtgacag cgccgncgtc ttcgcttga 360
 agggtgagac cctccaggag tactggtggt gcaccgagcg cgcccttgac tggggccccc 420

gcggcgggccc cgacctnatt gtcgacgacn gaagcnacgc cacctctnat cacgaaggcc 480
tnaangncca gggctctatn annaagaccg gggaactccc 520

<210> 3015
<211> 344
<212> nucleic acid
<213> Glycine max
<400> 3015

gttatttctc agcgcgtaaa gcatggcttt gttggtggag aaaaccacga gtggtcgcga 60
gtacaaggtc aaggaccttt cccaggccga cttcggccgc ctcgagatcg agctggccga 120
ggttgagatg cccggcctca tggcctgtcg naccgagttc ggcccntccc agccttcaag 180
ggggcccga tcaccggctc cctccacatg accatccage ggcgccgttc tcattgagac 240
cctcaccgcc cttggcgccg aggtccgctg gtgctcctgc aacatcttct ccanccagga 300
ccacgccgcc gccgtanttc ncgcgacagt gngccgtct tcgc 344

<210> 3016
<211> 528
<212> nucleic acid
<213> Glycine max
<400> 3016

ggnnnaagtt tttgnngggg ggggnannan gtntntntntnt nntntntntnt nntntntntnt 60
nnntntntntnt nntntntntnt nntntntntnt nntntntntnt nntntntntnt cganctggcc 120
cgaggtnag atggcgatcc tcatggcggtg gcggaccgag ttcggcccat ctcanccott 180
gaagggggcc cgcathnacc gctccttca catgaccatt canaccgntg tctcatoga 240
naccctnacc gctctcggcg ccgaggttcg ctggtgctcc tgcaacatct tctccactca 300
ggaccacgcc gncgncgcca tcgcccgtga cagcgccgnc gtcttcgcct ggaagggtga 360
gaccctncan gactactggt ggtncaccga gcgcgccctt gactggggcc cccgcgnggc 420
cccgacctta tngtcnaccg accgaaggca accccacctt cttatcaacn aggcggtnaa 480
ggccaaggag ctctatnaag aagaccgggg aactccccna ccctaact 528

<210> 3017
<211> 347
<212> nucleic acid

<213> Glycine max

<400> 3017

cgggtccct tcacatgacc atccagaccg ctgtcctcat cgagaccctn caccgctctc 60
ggcgccgagg ttcgctggtg ctctgcaac atctttctcca ctcaggacca nnnnnnnnnn 120
nnnnnnnnnc gtgacagcgc cgccgtcttc gcctggaagg gtgagaccct ccaggagtac 180
tggtggtgca ccgagcgcgc cctcgactgg ggccccggcg gcggccccga cctncatcgt 240
cgacgacgga ggcgacgcca cctcctcat ccacganggc gtcaaggcgg aggagctcta 300
tgagaagacc ggggaattcc ccgaccctaa ttccantgac aacggcg 347

<210> 3018

<211> 332

<212> nucleic acid

<213> Glycine max

<400> 3018

gtcgcangca cgcntacgtn agctcggaat tcggctcgag caggttctga ccttgaggga 60
tgttgtttct gaggctgata tctttgtcac caccaccggt aacaaggaca tcatcatggt 120
tgaccacatg aggaaaatga agaacaatgc cattgtttgc aacattggtc actttgacaa 180
tgagatcgac atgcttgggc tggagaacta ccccggcgtg aagcgcacat caatcaagcc 240
ccaaactgac agatgggtct tccctgagac caacaccggt atcattgtct tggtgaggg 300
tcgattgatg aacttgggat gcgccactgg ac 332

<210> 3019

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 3019

ngtngangca cgcgtacgta agctcggaat tcggctcgag gtcaaggacc tntcccaggc 60
cgacttcggc cgcctcgaga tcgagctggc cgaggttgag atgcccggcc tcatggcctg 120
tcggacngag ttcgggccct ccagccctt caagggggcc cgcacacccg gctccctcca 180
catgaccatc cagngcgccg ttctcattga gaccctcacc gcccttggcg ccgaggtccg 240
ctggtgctcc tgcaacatct tctccacca ggaccacgcc gccgcgcta ttgcccgca 300

cagtgcc

307

<210> 3020
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 3020

cgctgcntac gtacgtaagc tcggaattcg gctcgagccg agttcgggcc ctcccagccc 60
ttcaaggggg cccgcatcac cggctccctc cacatgacca tccagaccgc cgttctcatt 120
gagaccctca ccgcccttgg cgccgaggtc cgctggtgct cctgcaacat cttctccacc 180
caggaccacg ccgccgccgc tattgcccgc gacagtgcgc cgtcttctgc ctggaagggt 240
gagaccctcc aggagtactg gtggtgcacc gagegcggcc tcgactgggg ccccggtg 298

<210> 3021
<211> 339
<212> nucleic acid
<213> Glycine max

<400> 3021

ctttctctag tctgttatt tctcagcgcg taaagcatgg ctttgttggg ggagaaaacc 60
acgagtggtc gcgantacaa ggtcaaggac tttcccaggc cgacttcggc cgctcgcaga 120
tcgagctggc cgaggttgan atgcccggcc tcatggcctg tcggaccgag ttcgccccct 180
cccagccctt caagggggcn cgcatacccg gntccctcca catnaccatc cagaccgccg 240
ttctcattga gacctcacc gcccttggcg ccgaggtncg ctggtgctcc tgcaacatct 300
tctccacca ggaccacgcc gngccgctat tgtcgcgaa 339

<210> 3022
<211> 275
<212> nucleic acid
<213> Glycine max

<400> 3022

caccactggg gttaagaggc tatatcagat gcaggcgaat gggactctac tcttccctgc 60
tattaatgtc aatgactctg ttaccaagag caagtttgac aacttgtaag ggtgccgtca 120
ctctctccct gatggtctga tgagggtac tgatgtgatg attgctggaa aggtggctgt 180

tgtggctgga tatggtgatg ttggcaaggg ttgtgctgct gcattgaagc aggctggtgc 240
tcgtgtcatc gtgactgaga ttgaccccat ttgtg 275

<210> 3023
<211> 320
<212> nucleic acid
<213> Glycine max

<400> 3023

cnntncacgc gtacgtnagc tcggaattcg gctcgaggag gaaaccacca ctggtgttaa 60
gaggctatat cagatgcagg cgaatgggac tctactcttc cctgctatta atgtcaatga 120
ctctgttacc aagancaagt ttgacaactt gtacgggtgc cgtcactctc tccctgatgg 180
tctgatgagg gctactgatg tgatgattgc tggaaagggtg gctgttgtgg ctggatatgg 240
tgatgttggc aagggttgtg ctgctgcatt gaagcaggct ggtgctcgtg tcatcgtgac 300
tgagattgac cccattgtgc 320

<210> 3024
<211> 306
<212> nucleic acid
<213> Glycine max

<400> 3024

atgtcgcggtt cacgcgtacg taagctcgga attcggctcg aggggtggaga aaaccacgag 60
tggtcgcgag tacaaggcca aggacctttc ccaggccgac ttcggccgcc tcgagatcga 120
gctggccgag gttgagatgc ccggcctcat ggctgtcgg accgagttcg gcccctccca 180
gcccttcaag ggggcccga tcaccggctc cctccacatg accatccaga ncgccgttct 240
cattgagacc ctcaccgccc ttggcgccga ggtccgctgg tgctcctgca acatcttctc 300
caccca 306

<210> 3025
<211> 518
<212> nucleic acid
<213> Glycine max

<400> 3025

agtttntntt nngggggggg gggggnaang agtancgctn agctatgacg tcgcatgcac 60

gcgtacgtaa gctcggaatt cggctcgagg agagatctat ctatctatca agatggcggt 120
 gttgggttgag aaaacaagca gtggaagga gtacaaggtg aaggacatga cgcaagccga 180
 ttctggaaga ttggaatcg agctggcgga ggttgaaatg cccggcctca tgtcctcccg 240
 caccgagttc ggccccctctc aacccttcaa gggcgctagg atcaccggct ccctncacat 300
 gaccatncaa accgncgtcc tcatcgagac cctnaccggc ctngggcgccg aggtcccgt 360
 ggtgctnctg caacatcttc ttcanccaag accatgccgg cgcgcatcgc cgggacagcg 420
 ccttctgtctt cgcttgaaa ggtgagaccc ttcaggaatc tgggtggtgca ccgagcgcg 480
 ccttgactgg ggccccngcg gcggcccgat ctnattgt 518

<210> 3026
 <211> 338
 <212> nucleic acid
 <213> Glycine max
 <400> 3026

ctccaaggtc agaacctgaa ggccttccat gagagcctga tatctttgtc accaccaccg 60
 gtaacaagga catcatcatg gttgaccaca tgaggaaaat gaagaacaat gccattgttt 120
 gcaacattgg tcactttgac aatgagatcg acatgcttgg gctggagaac taccocggcg 180
 tgaagcgcat caccatcaag ccccaaactg acagatgggt ctnccctgag accaacaccg 240
 gtatcattgt cttggctgag ggtcgattga tgacttggga tgcgccatgg acaccccagt 300
 tttgtgatgt cctgtcctt caccaacang tcattgtc 338

<210> 3027
 <211> 286
 <212> nucleic acid
 <213> Glycine max
 <400> 3027

gtaccaatgt cggcatcatt gtcttggccg agggctggtt gatgaacttg ggatgcgcca 60
 caggacaccc tagttttgtg atgtcctgct ccttcaccaa ccaggtcatt gctcagcttg 120
 agttgtggaa ggagaagagt accggcaagt acgagaagaa agtttacgtt ttgcccgaagc 180
 accttgatga caaggtggct gcacttcacc ttggcaaact tggagctaag ctcaccaagc 240
 ttagcccggc ccaggctgat tacatcagtg tgcctgttga gggtec 286

<400> 3028

<210>	3029
<211>	312
<212>	nucleic acid
<213>	Glycine max

<400> 3029

<210>	3030
<211>	280
<212>	nucleic acid
<213>	Glycine max

<400> 3030

1066

tccacatgac catccagacc gccgtttctca ttgagaccct caccgccctt ggcgccgagg 240
tccgctggtg ctcttgcaac atcttctcca cccaggacca 280

<210> 3031
<211> 324
<212> nucleic acid
<213> Glycine max

<400> 3031

ngangcacgc gtacgtaagt cggaattcgg ctcgagtgtt atttctcagc gcgtaaagca 60
tgggctttgt tgggtggagaa aaccacgagt ggtcgcgagt acaagggtcaa ggacctttcc 120
caggccgact tcggccgcct cgagatcgag ctggccgagg ttgagatgcc cggcctcatg 180
gcctgtcgga ccgagttcgg cncctcccag cccttcaagg gggcccgcac caccggctcc 240
ctccacatga ccatccagac cgccgttctc attgagacc tcaccgccct tggcgccgag 300
ntccgctggt gctcctgcaa catc 324

<210> 3032
<211> 303
<212> nucleic acid
<213> Glycine max

<400> 3032

acgtcgang caccgctaag tnagctcgga attcggctcg aggccectcc cagcccttca 60
agggggcccg catcaccggc tccctccaca tgaccatcca gancgcgct tctcattgag 120
accctcaccg cccttggcgc cgaggtccgc tgggtgctct gcaacatctt ctccaccag 180
gaccacgcgc ccgcgctat tgcccgcgac agtgccgcgc tcttcgcctg gaagggtag 240
accctccagg agtactggtg gtgcaccgag cgcgccctcg actggggccc cgggtggtgga 300
ccc 303

<210> 3033
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 3033

gtnnatncac gcgtacgtaa gtcggaatt cggctcgagg ctggtgctcc tgcaacatct 60

tctccaccca ggaccacgcc gccgccgcta ttgcccgcga cagtgccgcc gtcttcgcct 120
 ggaaggggtga gaccctccag gactactggt ggtgcaccga gcgcgccctc gactggggcc 180
 ccggtggtgg acccgacctc atcgtcgacg acggtggtga cgctaccctt ctcatccacg 240
 aaggcgtcaa ggccgaggag ctctatgaga agaccggcga actccccgac cccaactcca 300
 ccgacaac 308

<210> 3034
 <211> 294
 <212> nucleic acid
 <213> Glycine max
 <400> 3034

3034

gtcgcangca cgcgtacgta agctcggaat tcggctcgag caccggtaac aaggacatca 60
 tcatgggtga ccacatgagg aaaatgaaga acaatgccat tgtttgcaac attggtcact 120
 ttgacaatga gatcgacatg cttgggctgg agaactaccc cggcgtgaag cgcatcacca 180
 tcaagcccca aactgacaga tgggtcttcc ctgagaccaa caccggatatc attgtcttgg 240
 ctgagggtcg attgatgaac ttgggatgcg ccaactggaca cccagtttt gtga 294

<210> 3035
 <211> 332
 <212> nucleic acid
 <213> Glycine max
 <400> 3035

ntacagtgcg angcacgcgt acgttagctc ggaattcggc tcgagctcta tcagatgcag 60
 gogaatggga ctcttctctt cctgctant aatgtcaatg actctgtcac caagancaag 120
 ttgacaact tgtatgggtg ccgtcactct ctccctgatg gtctcatgag ggctaccgat 180
 gttatgattg ctggaaaggt ggctgttggt gctggatatg gtgatgttgg caagggttgt 240
 gctgctgcaa tgaagcaggc tgggtctcgt gtcacgtga ccgagattga tcccatctgt 300
 gcccttcagg ctctcatgga agccttcagg tt 332

<210> 3036
 <211> 287
 <212> nucleic acid
 <213> Glycine max

<400> 3036

tcncatcgca tgcacgcgta cgtaagctcg gaattcggct cgagcgagta caaggtaag 60
gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgcnc 120
ggcctcatgg nctgtcggac ngagttcggc nctcccagc cttcaaggg ggcccgcatc 180
accggctccc tccacatgac catccaganc gccgtttctca ttgagaccct caccgccctt 240
ggcgccgagg tccgctggtg ctctgcaac atottctcca ccagga 287

<210> 3037

<211> 326

<212> nucleic acid

<213> Glycine max

<400> 3037

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ttctgaggaa accaccactg gtgttaagag gctatatcag atgcaggcga atgggactct 180
actcttctg ctattaatgt caatgactct gttaccaaga gcaagtttga caactgttac 240
gggtgccgtc actctctccc tgatggtctg atgagggcta ctgatgtgat gattgctgga 300
aagggtggctg ttgtggctgg atatgg 326

<210> 3038

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 3038

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tgtttgcaac attggtcact ttgacaatga gatcgacatg cttgggctgg agaactacco 120
cggcgtgaag cgcacaccca tcaagcccca aactgacaga tgggtcttcc ctgagaccaa 180
caccggtatc attgtcttgg ctgagggctg attgatgaac ttgggatgag ccactggaca 240
ccccagtttg tgatgtcctg ctcttcacc aaccaggta ttgtcagct tgagttgtgg 300
aaggag 306

<210> 3039

<211> 259
 <212> nucleic acid
 <213> Glycine max
 <400> 3039

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 aaggggtgaga ccctccagga gtactggtgg tgcaccgagc gcgccctcga ctgggggcccc 180
 ggtggtggac ccgacctcat cgtcgacgac ggtggtgacg ctacccttct catccacgaa 240
 gcgtcaaggc cgaggagct 259

<210> 3040
 <211> 306
 <212> nucleic acid
 <213> Glycine max
 <400> 3040

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 ggccgcctcg agatcgagct ggccgaggtt gagatgcccg gcctcatggc ctgtcggacc 180
 gagttcggcc cctcccagcc cttcaagggg gcccgcatca ccggctccct ccacatgacc 240
 atccagancg ccgtttctcat tgagaccctc accgcccttg gcgccgaggt ccgctggtgt 300
 cctgca 306

<210> 3041
 <211> 312
 <212> nucleic acid
 <213> Glycine max
 <400> 3041

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 gaaggagaag agtaccggca agtacgagaa gaaggtttac gttttgcca agcaccttga 180
 tgagaaggtg gctgcacttc acctgggcaa acttgngct aagctgacct agcttngcaa 240
 gtcccaggct gattacatca gtgtgcctgt tganggtcca tacaagcctg ctcaactacg 300

gtactaagtg at

312

<210> 3042
<211> 330
<212> nucleic acid
<213> Glycine max

<400> 3042

cnaagtctcn ngcacgcgta cgtaantcgg aattcggctc gaggctggtg ctcttgcaac 60
atcttctcca cccaggacca cgccgcccgc gctattgccc gcggacagtt nccgccgnct 120
tcgcctggaa ggggtgagacc ctccaggagt actggtggtg caccgagcgc gccctcgact 180
ggggccccgg tgggtgaccc gacctcatcg tcgacgacgg tgggtgacgt acccttctca 240
tccacgaagg cgtcaaggcc gaggagctct atgagaagac cggcgaactc cccgacccca 300
actccaccga caacgccgag tttcagatct 330

<210> 3043
<211> 314
<212> nucleic acid
<213> Glycine max

<400> 3043

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ggagttaaga ggctctatca gatgcaggcg aatgggactc ttctcttccc tgcattaat 120
gtcaatgact ctgtcaccaa gagcaagttt gacaacttgt atgggtgccg tcaactctctc 180
cctgatggtc tcatgagggc taccgatgtt atgattgctg gaaaggtggc tgttggtggt 240
ggatatgggtg atgttggcaa ggttggtgctg ctgcaatgaa gcaggctggt gctcgtgtca 300
tcgtgancga gatc 314

<210> 3044
<211> 312
<212> nucleic acid
<213> Glycine max

<400> 3044

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tattaatgtc aatgactctg tcaccaagag caagtttgac aacttgatg ggtgccgtca 120

ctctctccct gatggtctca tgagggctac cgatgttatg attgctggaa aggtggctgt 180
 tgtggctgga tatggtgatg ttggcaaggg ttgtgctgct gcaatgaagc aggctggtgc 240
 tcgtgtcatc gtgaccgaga ttgatcccat ctgtgccctt caggctctca tggaaggcct 300
 caggttctga cc 312

<210> 3045
 <211> 307
 <212> nucleic acid
 <213> Glycine max
 <400> 3045

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 ctctcatggn aggccttcag gntctgacct tggaggatgt tgtttctgan gctgatctct 180
 ttgtcaccac cacoggtaac aaggacatca tcatgggtga ccacatgagg aaaatgaaga 240
 acaatgccat tgtttgcnac attggctcact ntgacaatga gatcgacatg cttgggctgg 300
 agaacta 307

<210> 3046
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 3046

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 cagttttgtg atgtctgct ccttcaccaa ccaggctatt gtcagcttg agttgtggaa 120
 ggagaagagt accggcaagt acgagaagaa ggtttacgtt ttgcccaagc accttgatga 180
 gaaggtggct gcacttcacc tgggcaaact tggngctaag ctgaccacgc ttagcaagtc 240
 ccaggctgat tacatcagtg tgctgttgga gggccatac aagcctgctc actacaggta 300
 ctaagtgatt gaga 314

<210> 3047
 <211> 316
 <212> nucleic acid
 <213> Glycine max

<400> 3047

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tttgttggtg gagaaaacca cgagtggctg cgagtacaag gtcaaggacc tttcccaggc 120
cgacttcggc cgcctcgaga tcgagctngc cgagggttgag atgcccggcc tcatggcctg 180
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atgaccatcc agaccgccgt tctcattgag accctcaccg cccttggcgc cgagggtccg 300
tggtgctctg caacat 316

<210> 3048

<211> 259

<212> nucleic acid

<213> Glycine max

<400> 3048

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tgaccacatg aggaaaatga agaacaatgc cattgtttgc aacattgggc actttgacaa 120
tgagatcgac atgcttgggc tggagaacta ccccggcgtg angcgcatca ccatcaagcc 180
ccaaactgac agatgggtct tccctgagac caacaccggc atcattgtct tggtgaggg 240
tcgattgatg aacttgga 259

<210> 3049

<211> 346

<212> nucleic acid

<213> Glycine max

<400> 3049

agaacgnnaa nangtcgcat gcacgcgtac gtaagctcgg gaattcggct cgagcagtgc 60
cgccgtcttc ncctggaagg gtgagaccct ccaggagtac tgggtggtgca ccgagcgcgc 120
cctcgactga ngccccgggtg gtggaccgga cctcatcgtc gacgacgggtg gtgacgctac 180
ccttctcatc cacgaaggcg tcaaggccga ggagctctat gagaagaccg gcgaactccc 240
cgaccnaac tccaccgaca acgccgagtt tcagatcggtg cttaccatca tcagagatgg 300
gttgaagacc gatcccacca ggtaccgcaa agtgaaggag cgtctc 346

<210> 3050

<211> 319
 <212> nucleic acid
 <213> Glycine max

<400> 3050

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 tcagcgcgta aagcatggct ttgttggtgg agaaaaccac gagtggtcgc gagtacaagg 120
 tcaaggacct ttcccaggcc gacttcggcc gctcgagat cgagctggcc gaggttgaga 180
 tgcccggcct catggcctgt cggaaccgagt tcggcccctc ccagcccttc aagggggccc 240
 gcatcaccgg ctccctccac atgaccatcc agaccgccgt tctcattgag accctcaccg 300
 cccttggcgc cgaggtccg 319

3051
 298
 nucleic acid
 Glycine max
 3051

<210> 3051
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 3051

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 gtggagaaaa ccacgagtgg tcgcnagtac aaggtcaagg acctttccca ggccgacttc 120
 ggccgcctcg agatcgagct ggccgaggtt gagatgccg gcctcatggc ctgtcggacc 180
 gagttcggcc cctcccagcc cttcaagggg gcccgcatca ccggtccct ccacatgacc 240
 atccagaccg ccgttctcat tgagaccctc accgcncttg gcgcggangt ccgctggg 298

<210> 3052
 <211> 317
 <212> nucleic acid
 <213> Glycine max

<400> 3052

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 tcccaggccg acttcggccg cctcgagatc gagctggccg aggttgagat gcccggcctc 180
 atggcctgtn cggaccgagt tcggcccctc ccagcccttc aagggggccc gcatcaccgg 240
 ctccctccac atgaccatcc agaccgccgt tctcattgag accctcaccg cccttggcgc 300

cgagggtccgc tgggtgct

317

<210> 3053
<211> 311
<212> nucleic acid
<213> Glycine max

<400> 3053

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gttgggggttt ctgaggaaac caccactgga gttaagaggc tctatacaga tgcaggcgaa 120
tgggactctt ctcttcctg ctattaatgt caatgactct gtcaccaaga gcaagtttga 180
caacttgtat ggggtccgct actctctccc tgatgggtct atgagggtta ccgatgttat 240
gattgctgga aagggtggctg ttgtggctgg atatgggtgat gttggcaagg gttgtgctgc 300
tgcaatgaag c 311

<210> 3054
<211> 308
<212> nucleic acid
<213> Glycine max

<400> 3054

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catcatcatg gttgaccaca tgangaaaat gaagaacaan cgccattgtt tgcaacattg 120
gtcactttga caatgagatc gacatgctgg ggctggagaa ctaccccggc gtgaagcgca 180
tcaccatcaa gccccaaacc gacagatggg tcttccccga gaccaatgtc ggcatcattg 240
tcttggccga gggtcgtttg atgaacttgg gatgcgccac aggacaccct agttttgtga 300
tgtcctgc 308

<210> 3055
<211> 347
<212> nucleic acid
<213> Glycine max

<400> 3055

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gagtggctgc gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat 180
 cgagctggcc gaggttgaga tgcccggcct catggnctgt nggaccgagt tcggccccctc 240
 ccanccttc aagggggccc gcatcacccg ctccctccac atgaccatcc agancgccgt 300
 tctcattgag accctcaacg gcottggcgc cgangtccgc tggtgct 347

<210> 3056
 <211> 349
 <212> nucleic acid
 <213> Glycine max
 <400> 3056

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 ccacgaaggc gtcaaggccg aggagctcta tgagaagacc ggcgaactcc ccgaccccaa 120
 ctccaccgac aacgccgagt ttcagatcgt gcttaccatc atcagagatg ggttgaagac 180
 cgatcccacc aggtaccgca agatgaagga gcgtctcgtt ggggtttctg aggaaaccac 240
 cactggagtt aagaggctct atcagatgca ggcgaatggg actcttctct tccctgctat 300
 taatgtcaat gactctgtca ccaagagcga gttgacaatt gtatgggtg 349

<210> 3057
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 3057

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 cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac aaggtcaagg 120
 acctttccca ggctgacttc ggccgcctcg agatcgagct ggccgaggtc gagatgcccg 180
 gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg gcccgcatca 240
 ccggctccct ccacatgacc atccagaccg ccgttctcat tgagaccctc accgcccttg 300
 gcgcnnaggt ccgct 315

<210> 3058
 <211> 339
 <212> nucleic acid
 <213> Glycine max

<400> 3058

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cnngtacaag gtcaaggacc ttcccaggc cgacttcggc cgcctcgaga tcgagctggc 180
cgaggttgag atgcccggcc tcattggcctg tcggaccgag ttccggccct cccanccctt 240
caagggggcc cgcattcccg gctccctcca catgaccatc cagaccgccg ttctcattga 300
gaccctcacc gcccttggeg ccgaggtccg ctgggtgctc 339

<210> 3059

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 3059

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cggcgtgang cgcattacca tcaagcccca aaccgacaga tgggtcttcc ccgagaccaa 180
tgtcggcatc attgtcttgg ccgagggctg tttgatgaac ttgggatgcg ccacaggaca 240
ccctagtttt gtgatgtcct gctctcacca accaggtcat tgctcagctt gagttgtgga 300
a 301

<210> 3060

<211> 331

<212> nucleic acid

<213> Glycine max

<400> 3060

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aggtcaagga ctttcccg gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180
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cccgcattcac cggctccctc cacatgacca tccagaccgc cgttctcatt gagaccctca 300
ccgcccttgg ccgcgagtcc gtatgtcctg c 331

<210> 3061
 <211> 294
 <212> nucleic acid
 <213> Glycine max

 <400> 3061

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 ctgttgtggc tggatatggt gatgttggca agggttgtgc tgctgcattg aagcaggctg 180
 gtgctcgtgt catcgtgact gagattgacc ccatttgtgc ccttcaggct ctcattggaag 240
 gccttcagggt tctgacottg gaggatgttg tttctgaggc tgatatcttg tcac 294

<210> 3062
 <211> 291
 <212> nucleic acid
 <213> Glycine max

 <400> 3062

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 cactctctcc ctgatggtct catgagggct accgatgtta tgattgctgg aaagggtggct 180
 gttgtggctg gatatggtga tgttggcaag ggttgtgctg ctgcaatgaa gcaggctggt 240
 gctcgtgtca tcgtgaccga gattgatccc atctgtgccc ttcaggctct c 291

<210> 3063
 <211> 293
 <212> nucleic acid
 <213> Glycine max

 <400> 3063

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 tgactctgtc accaagagca agtttgacaa cttgtatggg tgccgtcact ctctccctga 180
 tgggtctcatg agggctaccg atgttatgat tgctggaaaag gtggtgttg tggtggata 240
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<400> 3064

[illegible]

<400> 3065

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1980	12	M
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1982	14	M
1983	15	F
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1985	17	F
1986	18	M
1987	19	F
1988	20	M
1989	21	F
1990	22	M
1991	23	F
1992	24	M
1993	25	F
1994	26	M
1995	27	F
1996	28	M
1997	29	F
1998	30	M
1999	31	F
2000	32	M
2001	33	F
2002	34	M
2003	35	F
2004	36	M
2005	37	F
2006	38	M
2007	39	F
2008	40	M
2009	41	F
2010	42	M
2011	43	F
2012	44	M
2013	45	F
2014	46	M
2015	47	F
2016	48	M
2017	49	F
2018	50	M
2019	51	F
2020	52	M
2021	53	F
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2025	57	F
2026	58	M
2027	59	F
2028	60	M
2029	61	F
2030	62	M
2031	63	F
2032	64	M
2033	65	F
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2035	67	F
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2044	76	M
2045	77	F
2046	78	M
2047	79	F
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2055	87	F
2056	88	M
2057	89	F
2058	90	M
2059	91	F
2060	92	M
2061	93	F
2062	94	M
2063	95	F
2064	96	M
2065	97	F
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2067	99	F
2068	100	M

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catggcctgt cgcccgagtt cgccccctcc cagcccttca aggggggccg catcaccggc 240
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 gaggtccgct ggtgctctgc aacat 325

<210> 3067
 <211> 296
 <212> nucleic acid
 <213> Glycine max
 <400> 3067

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 caagagcaag ttgacaact tgtaogggtg ccgtcactct ctccctgatg gtctgatgag 240
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 <212> nucleic acid
 <213> Glycine max
 <400> 3068

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 tggt 304

<210> 3069
 <211> 314
 <212> nucleic acid
 <213> Glycine max
 <400> 3069

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aaggccgagg agctctatga gaagaccggc gaactccccg accccaactc caccgacaac 240
gccgagtttc agatcgtgct taccatcatt agagatgggt tgaagaccga tcccaccagg 300
taccgcaaga tgaa 314

<210> 3070
<211> 299
<212> nucleic acid
<213> Glycine max

<400> 3070

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acaccctagt tttgtgatgt cctgctcctt caccaaccag gtcattgctc agcttgagt 299

<210> 3071
<211> 302
<212> nucleic acid
<213> Glycine max

<400> 3071

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ca 302

<210> 3072
<211> 289
<212> nucleic acid
<213> Glycine max

<400> 3072

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gtcggcatca ttgtcttggc cgagggctcg ttgatgaact tgggatgcnc cacaggacac 120
cctagttttg tgatgtcctg ccccttcacc aaccagggtca ttgctcagct tgagttgtgg 180
aaggagaaga gtaccggcaa gtacgagaag aaagtttacg ttttgcccaa gcaccttgat 240
gagaagggtg ctgcncttca ccttggcaaa cttgngcta agctcacca 289

<210> 3073

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3073

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tggtgtttgt ggctggatat ggtgatgttg gcaagggttg tgctgcacca ttgaagcngg 180
ctggtgctcg tgtcatcgtg actgagattg accccatttg tgcccttcag gctctcttgg 240
aaggccttca ggttctgacc ttggaggatg ttgtttctga ggctga 286

<210> 3074

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 3074

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gttgggggtt ctgaggaaac caccactgga gttaagaggc tctatcagat gcaggcgaat 120
gggactcttc tcttccctgc tattaatgtc aatgactctg tcaccaagag caagtttgac 180
aacttgatg ggtgccgtca ctctctccct gatgggtctc tgagggctac cgatgttatg 240
attgctggaa aggtggctgt tgtggctgga tatgggtgatg ttggc 285

<210> 3075

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 3075

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atcatgggtg accacatgaa gaaaatgaag aacaatgcc a ttgtttgcaa cattggtcac 120
tttgacaatg agatcgacat gctggggctg gagaactacc ccggcgtgan gcgcattcac 180
catcaagccc caaaccgaca gatgggtctt ccccgagacc aatgtcggca tcattgtctg 240
ggccgagggg cgtttgatga antgggatgc gccacaggac accctagttt tgtgatgtcc 300

<210> 3076

<211> 264

<212> nucleic acid

<213> Glycine max

<400> 3076

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ttgccgcgna cagtgccgcc gtcttcnctt ggaagggatga gacctccag gactactggg 120
ggtgcaccga gcgcgccctg cgactggggc cccggtggtg gaccgcacct catcgctgan 180
nacggtggtg acgctaccct tctcatccag gaaggcgtca aggcgcgagga gctctatgag 240
aagaccggcg aactccccga ncct 264

<210> 3077

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 3077

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gtaaagcatg gctttgttgg tggagaaaac cacgagtggg gcgcgagtaca aggtcaanga 120
cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg agatgcccg 180
cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg ccgcgcatcac 240
cggctccctc cacatgacca tenantcaaa ngttctcatt gagaccctca ccgcccttgg 300
cgccgaggtc 310

<210> 3078

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 3078

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 ttctctagtc ctgttatttc tcagcgcgta aagcatggct ttgttggtgg agaaaaccac 120
 gagtggtcgc gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat 180
 cgagctggcc gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggcccctc 240
 ccagcccttc aagggggccc gcatcaccgg ctccctccac atgaccatcc agancgccgt 300
 totcattgag accctcaccg ccttt 325

<210> 3079

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 3079

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 aaccaccact ggagttaaga ggctctatca gatgcaggcg aatggnactc ttctcttccc 120
 tgctattaat gtcaatgact ctgtcaccan gagcaagttt gacaacttgt atgggngccg 180
 tcactctctc cctgatggtc tcatganngc taccgatntt atgattgctg gaaaggtggc 240
 tgttggtggn ggatatggtg atgttggnan gggttgtgct gctncaatnn agcaggtctg 300
 tgcctenc 307

<210> 3080

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 3080

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 gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 120
 gacctttccc aggcgcactt cggccgcctc gagatcgagc tggccgaggt tgagatgcc 180
 ggctcatgg cctgtcggac cgagttcggc cctcccagc ccttcaaggg ggcccgcac 240
 accggctccc tccacatgac catccaganc gccgtttctc ttgagacct caccgccctt 300

ggn

303

<210> 3081
<211> 293
<212> nucleic acid
<213> Glycine max

<400> 3081

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catggctttg ttggtggaga aaaccacgag tggtcgogag tacaaggta aggacctttc 120
ccaggccgac ttcgcccgcc tcgagatcga gctggccgag gttgagatgc ccggcctcat 180
ggcctgtcgg accgagttcg gccctccca gcccttcaag ggggcccgca tcaccggctc 240
cctccacatg accatccaga ccgcggttct cattgagacc ctcaccgccc ttg 293

<210> 3082
<211> 309
<212> nucleic acid
<213> Glycine max

<400> 3082

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agcgcgtaaa gcatggcttt gttggtggag aaaaccacga gtggtcgca gtacaaggta 120
aaggaccttt ccaggccga ctctggccgc ctcgagatcg agctggccga ggtcgagatg 180
cccggcctca tggcctgttc ggaccgagtt cggccctcc cagcccttca agggggcccg 240
catcacgggc tcctccaca tgaccatcca gaccgcggtt ctcattgaga ccctcaccgc 300
ccttggcgc 309

<210> 3083
<211> 295
<212> nucleic acid
<213> Glycine max

<400> 3083

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gcgtaaagca tggctttggt ggtggagaaa accacgagtg gtcgcgagta caaggtaag 120
gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgccc 180

ggcctcatgg cctgtcggac cgagttcggc ccctcccagc cttcaaggg ggcccgcac 240
accggctccc tccacatgac catccagacc gccgtttctca ttgagaccct caccg 295

<210> 3084
<211> 303
<212> nucleic acid
<213> Glycine max
<400> 3084

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ggacctttcc caggccgact tcggccgcct cgagatcgag ctggccgagg ttgagatgcc 180
cggcctcatg gcctgtcggg cagagttcgg ccctcccag cccttcaagg gggcccgcac 240
caccggctcc ctccacatga ccacccagan cncgtttctc attgagacc tcaccgccct 300
tgg 303

<210> 3085
<211> 293
<212> nucleic acid
<213> Glycine max
<400> 3085

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gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgcc 180
ggcctcatgg cctgtcggac cgagttcggc ccctcccagc cttcaaggg ggcccgcac 240
accggctccc tccacatgac catccagacc gccgtttctca ttgagaccct cac 293

<210> 3086
<211> 322
<212> nucleic acid
<213> Glycine max
<400> 3086

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tcgagccggt cctgttattt ctcagcgct aaagcatgnc tttgttggtg gagaaaacca 120

cgagtgggtcg cgagtacaag gtcaaggacc tttcccaggc cgacttcggc cgctcgcaga 180
 tcgagctggc cgagggttgag atgcccggcc tcatggcctg tcggaccgag ttcgggccct 240
 cccagccctt caaggggggccc cgcaccccg gctccctcca catgaccatc cagancgccg 300
 ttctcattga gacctcacc gc 322

<210> 3087
 <211> 299
 <212> nucleic acid
 <213> Glycine max
 <400> 3087

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 aaccagggtca ttgctcagct tgagttgtgg aaggagaaga gtaccggcaa gtacgagaag 120
 aaagtttacg ttttgcccaa gcaccttgat gagaagggtg ctgcacttca ccttggcaaa 180
 cttggagcta agctcaccaa gcttagcccg gccagggtg attacatcag tgtgcctggt 240
 gaggggtccat acaagcctgc tcattacagg tactaagtaa ttgagattat caacggaaa 299

<210> 3088
 <211> 321
 <212> nucleic acid
 <213> Glycine max
 <400> 3088

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 gngtgggtcg gagtacaagg tcaaggacct ttcccaggcc gaattcggcc gctcagat 180
 cgagctggcc gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggccctc 240
 ccagcccttc aagggggccc gcaccccg ctccctccac atgaccatcc agaccgccgt 300
 tctcattgag acctcaccg c 321

<210> 3089
 <211> 304
 <212> nucleic acid
 <213> Glycine max
 <400> 3089

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 ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cacgagtggg cgcgagtaca 120
 aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180
 agatgcccgg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240
 cccgcatcac cggtccctc cacatgacca tccagtcgc cgttctcatt gagaccctca 300
 ccgc 304

<210> 3090
 <211> 318
 <212> nucleic acid
 <213> Glycine max

<400> 3090
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 gtacaaggtc aaggaccttt cccaggccga cttcggccgc ctcgagatcg agctggccga 180
 gggttgagatg cccggcctca tggcctgtcg gaccgagttc ggcccctccc agcccttcaa 240
 gggggcccgc atcaccggct cctccacat gaccatccag accgcccgttc tcattgagac 300
 cctcaccct tggcgccg 318

<210> 3091
 <211> 279
 <212> nucleic acid
 <213> Glycine max

<400> 3091
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 agtacaaggt caaggacctt tcccaggccg acttcggccg cctcgagatc gagctggccg 120
 aggttgagat gcccgccctc atggcctgtc ggaccgagtt cggcccctcc cagcccttca 180
 agggggcccg catcaccggc tccctccaca tgaccatcca gaccgcccgtt ctcattgaga 240
 ccctcaccgc ccttggcggn gacnncgggn nctnaaaaa 279

<210> 3092
 <211> 301
 <212> nucleic acid

<213> Glycine max

<400> 3092

gcacgcgtac gtaagctcgg aattcggctc gagntttctct agtcctgtta tttctcagcg 60
cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcnagtac aaggtcaagg 120
acctttccca ggccgaacttc ngccgcctcg agatcgagct ggccgagggt gagatgcccg 180
gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg gcccgcatca 240
ccggctccct ccanatgacc atccagaccg ccgtttctcat tgagacnctc accgcccttg 300
g 301

<210> 3093

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 3093

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ttggtcaactt tgacaatgag atcgacatgc tggggctgga gaactacccc ggcgtagangc 120
gcateaccat caagccccaa accgacagat gggctctccc gagaccaatg tcggcatcat 180
tgtctggcgg agggctggtt gatgaacttg ggatgcgcca caggacaccc tagttttgtg 240
at 242

<210> 3094

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 3094

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ctcagcgctt aaagcatggc tttgttggtg gagaaaacca cgagtggctc cgagtacaag 120
gtcaaggacc tttcccaggc cgacttcggc cgctcagaga tcgagctggc cgaggctcag 180
atgcccgccc tcatggcctg tcggaccgag ttccggccct cccagccctt caagggggcc 240
cgcatcaccg gtcctccac atgaccatcc agaccgccgt tctcattgag accctcaccg 300
ccc 303

<400> 3095

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<400> 3096

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<400>	3097
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cccgccctca tggcctgtcg gaccgagttc ggccccctccc ancccttcaa gggggcccgc 240
 atcaccggct cctccacat gacntccag accgcggttc tcattgagac cctcanogcc 300
 cttggcgcc 309

<210> 3098
 <211> 272
 <212> nucleic acid
 <213> Glycine max
 <400> 3098

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 cgttgggggtt totgaggaaa ccaccactgg agttaagagg ctctatcaga tgcaggcgaa 120
 tgggactctt ctcttccctg ctattaatgt caatgactct gtcaccaaga gcaagtttga 180
 caacttgat gggtgccgct actctctccc tgatgggtctc atgagggcta ccgatgttat 240
 gattgctgga aaggtggctg ttgtggctgg at 272

<210> 3099
 <211> 339
 <212> nucleic acid
 <213> Glycine max
 <400> 3099

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 gantacaang tgcaagganc tttcccaggc cgacttcggc ngntcgcaga tcgagctggc 120
 cgaggttgag atgcccgccc tcatggcctg tcggncgcag ttcggcncct ccagccctt 180
 caagggggcn cgcataccg gntccctcca catgaccatc cagancccg ttctcatttg 240
 agatcctnat cgccttggn gccgnaggtc cgctgggtct cctgnaacat cgtctccatc 300
 caggaccacg ccncngccgc tattgcccg anagtgcg 339

<210> 3100
 <211> 262
 <212> nucleic acid
 <213> Glycine max
 <400> 3100

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cgcgagtaca aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg 120
 gccgaggttg agatgcccg cctcatggcc tgtcggaccg agttcggccc ctcccagccc 180
 ttcaaggggg ccgcacatcac cggtccctc cacatgacca tccagctccg ccgtttctcat 240
 tgagaccctc accgccttg gc 262

<210> 3101
 <211> 276
 <212> nucleic acid
 <213> Glycine max
 <400> 3101

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 gcaggttgtt gctcgtgtca tcgtgactga gattgacccc atttnggccc ttnaggttct 120
 catggaaggc cttcaggttc tgaccttga ggatgttgtt tctgaggctg atatctttgt 180
 caccaccaag ggtaacaagg acatcatcat ggttgaccac atgaagaaaa tgangancan 240
 tgccattgtt tgcaacattg gtcactttga caatga 276

<210> 3102
 <211> 296
 <212> nucleic acid
 <213> Glycine max
 <400> 3102

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 aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180
 agatgcccg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240
 ccgcacatcac cggtccctc cacatgacca tccagaccgc cgtttctatt gagacc 296

<210> 3103
 <211> 294
 <212> nucleic acid
 <213> Glycine max
 <400> 3103

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tttcacatga ccatccagac cgtgtgctct atcgagaccc tcaccgctct cggcgccgag 120
 gttcgtgtgt gtcctgcaa catcttctcc actcaggacc acgcccgcgc cggcatcgcc 180
 cgtgacagcg ccgcgctctt cgcctggaag ggtgagaccc tccaggagta ctggtngtgc 240
 accgagcncg cctcgactg gggccccggc ggcgccccg acctcatcgt cgac 294

<210> 3104
 <211> 291
 <212> nucleic acid
 <213> Glycine max
 <400> 3104

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 gacctttccc aggcgactt cggcncctc gagatcgagc tggccgaggt tgagatgccc 180
 ggctcatgg cctgtcggac cgagttcggc cctcccagc cttcaaggg ggcccgcac 240
 accggtccc tccacatgac catccagacc gccgttctca ttgagaccct c 291

<210> 3105
 <211> 311
 <212> nucleic acid
 <213> Glycine max
 <400> 3105

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 ggtggtgacg ctacccttct catccacgaa ggctcaagg ccgaggagct ctatgagaag 120
 accggcgaac tccccgaccc caactccacc gacaacgccg agtttcagat cgtgcttacc 180
 atcatcagag atggggtgaa gaccgatccc accagggtacc gcaagatgaa ggagcgtctc 240
 gttgggggtt ctgaggaaac caccactgga gttaagaggc tctatcagat gcaggcgatt 300
 gggccttttn t 311

<210> 3106
 <211> 301
 <212> nucleic acid
 <213> Glycine max
 <400> 3106

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 ccgacttcgg ccgcctcgag atcgagctgg ccgaggttga gatgcccggc ctcatggcct 180
 gttcggaccg agttcggccc ctcccagccc ttcaaggggg cccgcacac cggctccctc 240
 cacatgacca tccagaccgc cgtttctcatt gagaccctca ccgccttgnc gccgaggtcc 300
 g 301

<210> 3107
 <211> 291
 <212> nucleic acid
 <213> Glycine max

<400> 3107

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 actctgtcac caagagcaag ttgacaact tgtatgggtg ccgtcatctc tccctgatgg 180
 tctcatgagg gctaccgatg ttatgattgc tggaaagggtg gctgttgttg ctggatatgg 240
 tgatgttggc aagggttgtg ctgctgcaat gangcagggtg gtccccgttc a 291

<210> 3108
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 3108

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 tgacgctacc cttctcatcc acgaaggcgt caaggccgag gagctctatg agaaaccggc 180
 gaactccccg accccaactc caccgacaac gccgagattc agatcgtgct taccatcatc 240
 agagatgggt tgaagaccga tcccaccagg taccgcaaga tgaaggagcg tctcgttg 298

<210> 3109
 <211> 341
 <212> nucleic acid
 <213> Glycine max

<400> 3109

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acaatgccat tgtttgcaac attggctact ttgacaatga gatcgacatg ctggggctgg 120
agaactaccc cggcgtgaag cgcataacca tcaagcccca aacngacaga tgggtcttcc 180
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ccacaggaca ccctagtttt gtgatgtctg tnccttcacc aaccaggtea tgctcagttg 300
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<210> 3110

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 3110

ngtcgcatgc acgcgtacgt aagctcgga ttcggctcga gcggctcgag gcaacatctt 60
ctccaccag gaccangcgg ccgcgctat tgcccgcgac agtgcngccg tcttcgcctg 120
gaaggggtgan accctccagg agtactggtg gtgcaccgag cgcgccctcg actggggccc 180
cgggtggtgga ccgacctca tcgtcgacga cgggtggtgac gctacccttc tcatccacga 240
aggcgtcaag gccgaggagc tctatnagaa gaccggcga 279

<210> 3111

<211> 271

<212> nucleic acid

<213> Glycine max

<400> 3111

ccaccangta ccgcaagatg aaggagcgtc tcgttgggggt ttctgaggaa accaccactg 60
gtgttaanan gctatatcag atncaggcna atgggantct actcttccct gctattaatg 120
tcaatgactc tgttaccaag agcaagtttg acaacttgta cgggtgccgt caactctctc 180
ctgatggtct gatgagggct actgatgtga tgattgctgg aaaggtggct gttgnggcc 240
ggatanggtg atnttgga gggttngcn c 271

<210> 3112

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 3112

agtcgcangc acgcgtacgt aagctcggaa ttccggtoga gtctctagtc ctgttatttc 60
tcagcgcgta aagcatggct ttgttggtgg agaaaaccac gantggtcgc gagtacaagg 120
tcaaggacct ttcccaggcc gacttcggcn gcctcgagat cgagctggcc gaggttgaga 180
tgcccggcct catggcctgt cggaccgagt tcggcccctc ccagcccttc aagggggccc 240
gcatcaccgg ctcccctcac atgaacctcc agaccgccgt tctcattgag acc 293

<210> 3113

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 3113

nngtcgcang caccgcgtacg taagctcggg attcggctcg agtctttctc tagtcctgtt 60
atctctcagc gcgtaaagca tggctttgtt ggtggagaaa accacgagtn gtcgcgagta 120
caagggtcaag gacctttccc aggcgcactt cggccgcctc gagatcgagc tggccgaggt 180
tgagatgccg ggccctcatgg cctgtcggac cgagttcggc cctcccagc ccttcanggg 240
ggcccgcatc accggctccc tccacatgac catccagacc gccgtttctc ttgagaccct 300
n 301

<210> 3114

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 3114

tcgcangcac gcgtacgtaa gctcgggaatt cggctcgagt nggccgcctc gagatcgagc 60
tggccgaggt tgagatgccg ggccctcatgg cctgccggac cgagttcggc ccatctccag 120
cccttcaagg gggcccgcat caccggcncc cttcacatga ccatcnagac cgtgtctctc 180
atcgagaccc tcaccgctct cggcgccgag gttcgtgtgt gtcctgcaa catcttctcc 240
actcaggacc acgccgccgc cgccatcgcc cgtgacagcg ccg 283

<210> 3115

<211> 313
 <212> nucleic acid
 <213> Glycine max

 <400> 3115

 gtcgcangca cgcgtacgtn nagotcggaa ttcggtcga gctttctcta gtctgttat 60
 ttctcagcgc gtaaagcatg gctttgtngg tggagaaaac cagagtggt cgcgagtaca 120
 aggtcaagga cctttcccag gcgacttcg gccgcctcga gatcgagctg gccgaggttg 180
 anatgcccgg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240
 cccgcacac cggctccctc cacatgccat ccagaccgcc gttctcattg anaccctnac 300
 ngcccttggg cga 313

<210> 3116
 <211> 305
 <212> nucleic acid
 <213> Glycine max

 <400> 3116

 ncgtcgcag cgcggtacg tnagctcga attcggtcga agctctttct ctagtcctgt 60
 tttttctcag cgcgtaaagc atggccttgt tggcggagaa aaccacgagt ggtcgcgagt 120
 acaaggncna ggacctttcc caggccgact tcggccgcct cgagatcgag ctggccgagg 180
 ttgagatgcc cggcctcatg gctgtcga ccgagttcgg cccctcccag ccttcaagg 240
 gggcccgcat caccggtcc ctccacatga ccatccagan cgcggttctc attgagacct 300
 caccg 305

<210> 3117
 <211> 279
 <212> nucleic acid
 <213> Glycine max

 <400> 3117

 gtcgcangca cgcgtacgta agctcggaa tcggtcga gttatttctc agcgcgtaaa 60
 gcatggcttt gttggtggag aaaaccacga gtggtcga gtacaaggtc aaggaccttt 120
 cccaggccga cttcggccgc ctcgagatcg agctggccga ggttgagatg cccggcctca 180
 tggcctgtcg gaccgagttc ggcccctccc agcccttcaa gggggccgc atcaccggct 240

ccctccacat gaccatccag acagccgttc tcattgaga

279

<210> 3118
<211> 301
<212> nucleic acid
<213> Glycine max

<400> 3118

agtcgcangc acgcgtacgt aagctoggaa ttcggtcga gntagtcctg ttattttctca 60
gcgcgtaaag catggctttg ttggtggaga aaaccacgag tgggtgcgca gtacaaggtc 120
aaggaccttt cccaggccga cttcggcgc ctcgagatcg agctggccga ggttgagatg 180
cccggcctca tggcctgtcg gaccgagttc ggccccctcc agcccttcaa gggggcccgc 240
atcaccggct ccctccacat gaccatccag accgccgttc tcattgagac ctcaccgcc 300
c 301

<210> 3119
<211> 322
<212> nucleic acid
<213> Glycine max

<400> 3119

ngatgntgcn nncgccccgn agatcggaat tncgggctcg agctgtgacc tntcaggatc 60
tcanggnagg ccttcaggnt ctgaccttng aggatgttng ttctgaggct gatatcngtg 120
tcaccancca ncggtaacaa ggacatcatc atgggtgacc acatgangan aatgaagaac 180
aatgccattg tttgcaacat tggtcatttg acaatgagat cgacatgctt gggctggaga 240
actaccccgg cgtgaagcgc atcaccatca agccccaac tgacagatgg gtcttccctg 300
agaccaaacac cggatcatgt ct 322

<210> 3120
<211> 293
<212> nucleic acid
<213> Glycine max

<400> 3120

gtcgcangca cgcgtacgta agctcggaat tccggtcga ctctttctct agtcctgtta 60
tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120

aaggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgaggtt 180
gagatgnccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
gcccgcatca ccggtccct ccacatgacc atccagatcg ccgttctcat tga 293

<210> 3121
<211> 313
<212> nucleic acid
<213> Glycine max

<400> 3121

ttcatgcacg cgtaohtaag ctcggaattc ggctcgagga gagagagaga gatctatcta 60
tctatcaaga tggcggttgtt ggttgagaaa acaagcagtg gaagggagta caaggtgaag 120
gacatgacgc aagccgattt cggaagattg gaaatcgagc tggcggaggt tgaaatgccc 180
ggcctcatgt cctcccgac cgagttcggc ccctctcaac cttcaaggg cgctaggatc 240
accggtccc tccacatgac catccaaacc gccgtcctca tcgagaccct caccgccctc 300
ggcgccgagg tcc 313

<210> 3122
<211> 315
<212> nucleic acid
<213> Glycine max

<400> 3122

nngatgnacg cgtacgttag ctcggaattc ggctcgagct cttctcttag tccgtttatt 60
tctcagcgcg taaagcatgg ctttggttgg ggagaaaacc acgagtgggc gcgagtacaa 120
ggtcaaggac ctttcccagg cngacttcgg ccgcctcgag agcgagctgg ccgaggttga 180
gatgcccggc ctcatggcct gtcggaccga gttcggcccc tccagccct tcaagggggc 240
ccgcatcacc ggctccctcc antgaccatc cagttccgcc gttctcattg agaccctcac 300
cgcccttggc gccga 315

<210> 3123
<211> 297
<212> nucleic acid
<213> Glycine max

<400> 3123

aanagcatgc acgcgtacgt aagctcgga ttcggctcga gccactctct ttctctagtc 60
 ctgttatttc tcagcgcgta aagcatggct ttgttggtgg agaaaaccac gagtggtcgc 120
 gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat cgagctggcc 180
 gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggcccctc ccagcccttc 240
 aagggggccc gcatcaccgg ctccctccac atgaccatcc agaccgccgt tctcatt 297

<210> 3124
 <211> 290
 <212> nucleic acid
 <213> Glycine max
 <400> 3124

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctttctct agtcctgtta 60
 tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120
 aaggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgaggtt 180
 gagatgcccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
 gcccgcatca ccggtccct ccacatgacc atccagaccg ccgttctcat 290

<210> 3125
 <211> 273
 <212> nucleic acid
 <213> Glycine max
 <400> 3125

angtcgang cagcgtacg ttagctcgga attcggctcg agagaaaatg aagaacaatg 60
 ccattgtttt caacattggt cactttgaca atgagatcga catgctgggg ctgganaact 120
 accccggcgt gangcgcac accatcaagc cccaaaccga cagatggtct tccccgagac 180
 caatgtcggc atcattgtct tggccgaggg tcgtttgatg aacttgggat gcgccacagg 240
 acaccctagt tttntgatgn cctgctcctt cac 273

<210> 3126
 <211> 289
 <212> nucleic acid
 <213> Glycine max
 <400> 3126

ctnccgcangc ncgcgtacgt aagctcggaa ttcggctcga gctctttctc tagtcctggt 60
 atttctcagc gcgtaaagca tggctntggt ggtggagaaa accacgagtg gtcgcgngta 120
 caagggtcaag gacctttccc aggcgcactt cggcgcctc gagatcgagc tggccgaggt 180
 tgagatgcc ggctcatgg cctgtcggac cgagttcggc ccctcccagc cttcaagggt 240
 ggccgcgcatc accggtccc tccacatgac catccaganc gccgttctc 289

<210> 3127
 <211> 310
 <212> nucleic acid
 <213> Glycine max

<400> 3127

anncacgcgt acgtaagctc ngaattcggc tcgagctctt tctctagtc tgnatttctn 60
 cancgngtaa agcatggctt tnttggtgga gaaaaccacg agtngtcgcn agtacaatgt 120
 caaggacctt tcccaggcgc acttcggncg cctngagatc ganctggccg aggttganan 180
 gcacggcctc atggcctgtc ggaccgagtt cngcccctcc cancccttca agggggcccg 240
 catcaccggc tccctccaca tgaccatcca gaccggcggtt ctcatcgaga ccctcaccgn 300
 cttggcgccg 310

<210> 3128
 <211> 292
 <212> nucleic acid
 <213> Glycine max

<400> 3128

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga ggcgcctcgc actggggccc 60
 cgggtggtgga cccgacctca tcgtcgacga cgggtgtgac gctacccttc tcatccacga 120
 aggggtcaag gccgaggagc tctatgagaa gaccggcgaa ctccccgacc ccaactccac 180
 cgacaacgcc gagtttcaga tcgtgcttac catcatcaga gatgggttga agaccgatcc 240
 caccaggtac cgcaagatga aggagcgtct cgttgggggtt tctgaggaaa cc 292

<210> 3129
 <211> 299
 <212> nucleic acid
 <213> Glycine max

<400> 3129

nnngtatgca cgcgtacgta agctcggaat tcggctcgag ctttctctag tcctgttatt 60
tctcagcgcg taaagcatgg ctttggttgg ggagaaaacc acnctgggtg cgcgagtaca 120
aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180
agatgcccgg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240
cccgcatac cggctccctc cacatgacca tccagancgc cgttctcatt gagaccctc 299

<210> 3130

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 3130

aaaaaagnaa ngtcgcatgc acgcgtacgt aagctcgga ttcggctcga gtcctgttat 60
ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cacgagtggc ccgcgagtac 120
aaggtcaagg acctttccca gccgacttc gccgcctcga agatcgagct gccgaggttg 180
gagatgcccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
gcccgcatac cgggtccct ccacatgacc atccagaccg ccgttctcat tgagacctca 300
ccgccttggc gccgaggtcg ctnnn 325

<210> 3131

<211> 273

<212> nucleic acid

<213> Glycine max

<400> 3131

aacgcgtang taagctcgga attcggtcgc agntttctca gcgcgtaaag catggctttg 60
ttggtggaga aaaccacgag tggtcgcgag tacaaggta aggaccttcc ccaggccgac 120
ttcggccgcc tcgagatcga gctggccgag gttinatgc ncggcctcat ggctgtcgg 180
accgagttcg gnnctncca gcccttcaag ggggcncgca tcanoggntc cctccacatg 240
accatcnagn ccgccgttct cattgagacc etc 273

<210> 3132

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3132

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctttctct agtcctgtta 60
 tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120
 aagggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgagggtt 180
 gagangcccg gctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
 gcccgcatca ccggctccct ccacatgacc atccagaccg ccgttc 286

<210> 3133

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 3133

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctttctct agtcctgtta 60
 tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120
 aagggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgagggtt 180
 gagangcccg gctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
 gcccgcatca ccggctccct ccacatgacc atccagatcg ccgttctc 288

<210> 3134

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 3134

tcgcangcan gcacgcgtac gtaagctcgg aattcggctc gaggttatatt ctcagcgogt 60
 aaagcatggc tttgttggtg gagaaaacca cgagtggctg cgagtacaag gtcaaggacc 120
 tttcccaggc cgacttcggc cgctcggaga tcgagctggc cgaggttgag atgcccggcc 180
 tcatggcctg tcggaccgag ttccggcccct ccagcccctt caagggggcc cgcattcaccg 240
 gtcctctcca natgaccatn cagaccgccg tccctcattg agacctca 289

<210> 3135

<211> 289

<212> nucleic acid

<213> Glycine max
 <400> 3135
 nacgtcgcat gcaacgcgtac gtaagctcgg aattcggctc gaggtcgacg acggtggtga 60
 cgctaccott ctcatocacg aaggcgtcaa ggccgaggag ctctatgaga agaccggcga 120
 actccccgac cccaactcca ccgacaacgc cgagtttcag atcgtgctta ccatcatcag 180
 agatgggttg aagaccgatc ccaccaggta ccgcaagatg aaggagcgtc tcgttggggt 240
 ttctgaggaa accaccactg gagttaagag gctctatcag atgcaggcg 289

<210> 3136
 <211> 281
 <212> nucleic acid
 <213> Glycine max

3136
 <400> 3136
 tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc tagtcctgtt atttctcagc 60
 gognaaagca tggctttgtt ggtggagaaa accacgagtg gtcgagagta caaggtcaag 120
 gacctttccc agnccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgccc 180
 ggccatcatgg cctgtcggac cgagttcggc ccctcccagc ctttcaaggg ggcccgcac 240
 accggctccc tccacatgac catccatgac accgtttctca t 281

<210> 3137
 <211> 301
 <212> nucleic acid
 <213> Glycine max

<400> 3137
 gtcgctgcac gcgtacgtaa gctcggaatt cggctcgagc tagttttgtg atgtcctgct 60
 cnttcaccan nccaggatcat tgctcagctt gagttgtgga aggagaagag taccggcaag 120
 tacgagaaga aagtttacgt ttgcccgaag cacottgatg agaaggtggc tgcacttcac 180
 cttggcaaan ttggagctaa gtcaccaag cttagcccgg ccagggtga ttacatcagt 240
 gtgcctgttg agggccata caagcotgct cattacaggt actaagtaat tgagattatc 300
 a 301

<210> 3138

<211> 286
 <212> nucleic acid
 <213> Glycine max

<400> 3138

acgtcgcacg cagcggtacg taagctcgga attcggctcg agctttctct agtcctgtta 60
 tntctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120
 aaggtaagg acccttccca ggccgacttc ggccgctcg agatcgagct ggccgagggt 180
 gagatgcccg gctcatggn ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
 gnccgcatca ccggtccct ccacatgacc atccagancg ccgttn 286

<210> 3139
 <211> 270
 <212> nucleic acid
 <213> Glycine max

<400> 3139

ngatcgcntn aagtcgcang cagcggtacg tangctcggn aattcggctc gagntgacaa 60
 cttgtacnng tgccgtcact ctctccctga nggtctgatg agggctactg atgtgatgat 120
 tgctggaaag gtggctgttg tggctggata tggatgatgt ggcaaggggt gtgctgctgc 180
 attgaagcag gctggtgctc gtgtcatcgt gactgagatt gacccattt gtgccottca 240
 ggctctcatg gaaggcctca gttctgacct 270

<210> 3140
 <211> 298
 <212> nucleic acid
 <213> Glycine max

<400> 3140

ncgattgcac gcgtacgtna gctcggatt cggtctgagn ngagaccctc caggagtact 60
 ggtggtgcac cgagcgcgcc ctacgactgg ggccccggtg gtggaccgga cctcatcgtc 120
 gacgacggtg angacgtac cttctcatc cacgaaggcg tcaaggccga ggantctatg 180
 agacgaccgg cgaactcccc gaccccaact ccaccgacaa cgccgagttt cagatcgtgc 240
 ttaccatcat cnganatggg ttgaagaccg atcccaccag gtaccgcaag atgaagga 298

<210> 3141

<211> 334
 <212> nucleic acid
 <213> Glycine max

 <400> 3141

 gangacgcgt acgttagctc ggaattccgc tcgagctcga gccggggaaa ccaacactgg 60
 agttaagagg ctctatcaga tgcnaggcga atgggattcc tctcttccng ntaataaatg 120
 tcaatgactc ntgtcaccan gagcnagttt gacaacttgt atgggtgccg tncactctct 180
 ccctgatggt ctcatgaggg ctaccgatgt tatgattgct ggaaagggtg ctgttgtngc 240
 tggatatggt gatgttgga anggttgtgc tgctgcaatg naggaggctg gtgctcgtgt 300
 catcgtgnac gagattgac ccattctgtgc cctc 334

<210> 3142
 <211> 266
 <212> nucleic acid
 <213> Glycine max

 <400> 3142

 ctctctttct ctagtctgt tatttctcan cgcgtanagc atggctttgt tggaggagaa 60
 aaccacgagt ggtcgcgagt acaaggtcaa ggacctttcc caggccgact tcggccgcct 120
 cgagatcgag ctggccgagg ttgagatgcc cggcctcatg gcctgtcgga ccgagttcgg 180
 cccctcccag ccttcaagg gggcccgcat caccggtcc ctccacatna cnaanaaatn 240
 ncnaantctc attgagacc tcancg 266

<210> 3143
 <211> 288
 <212> nucleic acid
 <213> Glycine max

 <400> 3143

 ntgcncnngt acgtaagctc ggaattccgc tcgagctctt tctctagtcc tgttatttct 60
 cagcgcgtaa agcatggctt tgttggtgga gaaaaccacg agtggtcgcg agtacaaggt 120
 caaggacctt tcccaggccg acttcggccg cctcgagatc gagctggccg aggttgagat 180
 gcccggcctc atggcctgtc ggaccgagtt cggccctcc cagcccttca agggggcccg 240
 catcaccggc tcctccaca tgaccatcca ganncgccgt tctcattg 288

<210> 3144
 <211> 308
 <212> nucleic acid
 <213> Glycine max

 <400> 3144

 ncacgcgtac gtnagctcgg aattcggctc gagcctcgac gacggtggtg acgctaccct 60
 tctcatccac gaaggcggtca aggccgagga gctctatgag aagaccggcg aactccccga 120
 ccccaactcc accgacaacg ccgagtttca gatcgtgctt ancatcatca gagatggggtt 180
 gaagaccgat cccaccaggt ancgcaagat gaaggagcgt ctcggtggggg tttctgagga 240
 aaccaccatt ggagttaaga ggntctatca gatgcaggcg aatgggatct tctcttcctt 300
 gctattaa 308

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 1179
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 1188
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 1192
 1193
 1194
 1195
 1196
 1197
 1198
 1199
 1200

<210> 3145
 <211> 279
 <212> nucleic acid
 <213> Glycine max

 <400> 3145

 nntcgcacgc acgcgtacgt aagctcggaa ttcggctcga gctctttctc tagtcctggt 60
 atttctcagc gcgtaaagca tggctttggt ggtggagaaa accacgagtg gtcgcgagta 120
 caagggtcaag gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt 180
 tgagatgcc ggctcatgg cctgtcggac cgagttcggc cctcccagc ctttcaaggg 240
 ggcccgcatc accggtccc tccacatgac catccagac 279

<210> 3146
 <211> 296
 <212> nucleic acid
 <213> Glycine max

 <400> 3146

 cgtcgcangc acgcgtacgt nagctcggaa ttcggctcga gngcgacgcc accctcctca 60
 tccacgaggg cgtcaaggcc gaggagctct atgagaagac cggggaactc cccgacccta 120
 actccactga caacgccgag ntccagatcg tgcttaccat catcagagat gggttgaaga 180
 ccgatccac caggtaccgc aagatgaagg agcgtctcgt tggggtttct gaggaacca 240

ccactggtgt taagaggcta tatcagatgc aggcgaatgg gactctactc ttccct 296

<210> 3147
 <211> 287
 <212> nucleic acid
 <213> Glycine max

<400> 3147

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag cgtaaagcat ggctttgttg 60

gtggagaaaa ccacgagtgg tcgcgagtta caaggtcaag gacctttccc aggccgactt 120

cgcccgcttc gagatcgagc tggccgaggt tgagatgccc ggcctcaggc ctgttcggac 180

cgagttcggc cctcccagc cttcaaggg ggccgcctc accggtccc tccacatgac 240

catccagacc gccgtttctc tgagaccctc accgcccttg gcgccga 287

3148
 275
 nucleic acid
 Glycine max
 3148

<210> 3148
 <211> 275
 <212> nucleic acid
 <213> Glycine max

<400> 3148

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctttctct agtcctgtta 60

tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120

aaggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgaggtt 180

gagatgcccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240

gcccgcctca ccggctccct ccacatgacc atcca 275

<210> 3149
 <211> 239
 <212> nucleic acid
 <213> Glycine max

<400> 3149

atcgtagcta agctcggaat tcggctcgag cttaccaaac caggtcattg ctcagnttga 60

gttgtggatg gagaagagta ccggcaagta cgagaagaag gtttacgttt tgcccaagca 120

ccttgatgag aaggtggctg cacttcacct gggcaaactt ggngctaagc tgaccagct 180

tagcaagtcc caggctgatt acatcagtgt gcctgttgag ggtccatata agcctgctc 239

<210> 3150
 <211> 270
 <212> nucleic acid
 <213> Glycine max

 <400> 3150

 gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctttctctag tcctgttatt 60
 tctcagcgcg taaagcatgg ctttggttgg ggagaaaacc acgagtggc gcgagtacaa 120
 ggtcaaggac ctttcccagg ccgacttcgg ccgcctcgag atcgagctgg ccgagggtga 180
 gatgcccggc ctcatggcct gtcggaccga gttcggcccc tcccagccct tcaagggggc 240
 ccgcacacc ggctccctcc acatgaccat 270

<210> 3151
 <211> 290
 <212> nucleic acid
 <213> Glycine max

 <400> 3151

 gtcgcangca cgcgtacgta agctcgggaa ttccggctcga ggtaaagcat ggctttgttg 60
 gtggagaaaa ccacgagtgg tcgcgagtac aaggtaagg acctttccca ggccgacttc 120
 ggccgcctcg agatcgagct ggccgagggt gagatgcccg gcctcatggc ctgtcggacc 180
 gagttcggcc cctcccagcc cttcaagggg gcccgcatca ccggtccct ccacatnaca 240
 nnnnacngaa aaatgctcat tgagaccctt caccgccnnt gggggcgngg 290

<210> 3152
 <211> 310
 <212> nucleic acid
 <213> Glycine max

 <400> 3152

 agcannagnt cnngangcgt acgtaagtcg ganttcggct cgagntcttt ctctagtent 60
 gttatttctc agcggcgtaa agcatgggtt tgttgatgga gnaaaccaac gagtcgttng 120
 cgagtacaag gtcaaggacc tttcccaggc cgacttcggc cgctcgaga tcgagctggc 180
 cgagggttgag atgcccggcc tcatggcctg tcggancgag ttccggccct cccagccctt 240
 caagggggcc cgcacaccg gctccctcca natgaccatc cagaccgcg ttctcattga 300

gagctcacgc 310

<210> 3153
<211> 277
<212> nucleic acid
<213> Glycine max

<400> 3153

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctagtccctgt tatttctcag 60
cgcgtaaagc atggctttgt tgggtggagaa aaccacnagt ggtacgcgag tacaagggtca 120
aggacctttc ccaggccgac ttcggccgcc tcgagatcga gctggccgag gttgagatgc 180
ccggcctcat ggctgtcggt accgagttcg gccctccca gcccttcaag ggggcccga 240
tcaccggctc cctccacatg accatccaga ccgccgt 277

gagctcacgc
gtcgcangca
cgcgtaaagc
aggacctttc
ccggcctcat
tcaccggctc
cctccacatg
accatccaga
ccgccgt

<210> 3154
<211> 298
<212> nucleic acid
<213> Glycine max

<400> 3154

gcangcacgc gtacgtaagc tcggaattcg gctcgagctc tctttctcta gtcctgttat 60
ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cagcagtggt cgcgagtaca 120
aggtaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgagggtg 180
agatgcccgg cctcatggcc tntcggaccg agttnggcc cgnccagccc gtnaaggggg 240
cccgcacnc cggcgctcgc nacaggatca nccagaccgc cgttctcagt ganaccgc 298

<210> 3155
<211> 318
<212> nucleic acid
<213> Glycine max

<400> 3155

ngtcgcatgc acgcgtacgt aagctcgga atnnngctc gagctcgagc cgctncgagc 60
cgctccactc tctttctcta gtcctgttat ttctcagcgc gtaaancatg gctnanttgg 120
tggagaaaac cagcagtggt cgcnagtaca aggtcaagga cctttcccag gccgacttcg 180
gcgcctcgag atcgagctgg ccngggtga gatgcccgc ctcattggcct gtncggaccg 240

agttcgcccc ctcccagccc ttcaaggggg cccgcatcac cggctccctc cacatgacca 300
tccagaccgc cgtttctca 318

<210> 3156
<211> 318
<212> nucleic acid
<213> Glycine max

<400> 3156

cactctcttt ctctagtcct gttattttctc agcgcgtaaa gcatggcttt gttggtggag 60
aaaaccacga gtggtcgcga gtacaaggtc aaggaccttt ccagggccga cttcgggccgc 120
ctcgagatcg agctggccga ggttgagatc ccggcctcat ggctgtngg accgagttcg 180
gcccctcccag cctcaaggg ggcccgcac accggctccc tccacatgac catccagacc 240
gccgtttctca ttgagacctc acngccttgg gccgagtcgc ttggtgtctt gaaanatttc 300
tcaaccaagg acaagcng 318

<210> 3157
<211> 292
<212> nucleic acid
<213> Glycine max

<400> 3157

gttgacacgc tacgtaagct cggaattcgc ctcgagctct ctttctctag tctgtttatt 60
tctcngcgcg taaagcatgg ctttgttggg ggagaaaacc angagtgggc gcgagtacaa 120
ggtcaaggac gtttcccagg ccgacttcgc ccgcctcgag atcgagctgg ccgngggtga 180
gatgcccggc ctcatggcct gtcggaccga gttcggcccc tcccagccct tcaagggggc 240
ccgcatcacc ggctccctcc acatnacann cgacngcanc gttctcattg an 292

<210> 3158
<211> 278
<212> nucleic acid
<213> Glycine max

<400> 3158

tcgcangcac gcgtacgtna gtcggaatt cggctcgagn tttctctagt cctgttattt 60
ctcagcgcgt aaagcatggc tttgttgggt gagaaaacca cgagtggctg cgagtacaag 120

gtcaaggacc tttcccaggc cgacttcggc cgctcgcaga tcgagctggc cgaggttgag 180
 atgcccggcc tcatggcctg tcggaccgag ttcgggccct cccagccctt caagggggcc 240
 cgcatcaccg gctccctcca catggaccat ccagaccg 278

<210> 3159
 <211> 332
 <212> nucleic acid
 <213> Glycine max

<400> 3159

acaccncct accacgccaan cgnaagctcg gaattnggct cgagattcac caaccaggt 60
 cattgctcag ttgagttgtg gaaggagnag agtaccggca agtacgagaa gaaggtttac 120
 gttttgcnc aagcacttga tgagaagggtg gctgcactta acctgggcaa acttgnagct 180
 aagctgaccc agcttagcaa gtnccaggnt gattacatca gtgtgcctgt tgaggggtcca 240
 tacaagcctg ctcantacag gtacnnnctn atnnngatga tcaactgnaa agtgagtgag 300
 ggaaagacaa aaatgggttt tatnaatngg at 332

<210> 3160
 <211> 288
 <212> nucleic acid
 <213> Glycine max

<400> 3160

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc tctttctcta gtcctgttat 60
 ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cacgagtggc cgcgagtaca 120
 aggtcaagga cttttccag gccgacttcg gccgcctcga gatcgagtgg ccgaggttga 180
 gatgcccggc ctcatggcct ntcggaccga gttcgggccc tcccagccct tcaagggggc 240
 ccgcatcacc ggctccctcc acatgaccat ccagnngccg ttctcatt 288

<210> 3161
 <211> 282
 <212> nucleic acid
 <213> Glycine max

<400> 3161

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc tctttctcta gtcctgttat 60

ttctcagcgc gtaaagcatg gctttgttgg tggagataac cacnctggt ccncgagtac 120
aagggtcaagg acctttccca ngccgacttc ggccgcctcg agatcgagct ggccgaggtt 180
gagatgcncg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240
gcccgcatca ccggnctcct ccacatganc atccagaccg cc 282

<210> 3162
<211> 318
<212> nucleic acid
<213> Glycine max
<400> 3162

cgcacgncgc gaacnnnagc ncgcgaattc ggntcgagng ngcccnccgac tggggccccg 60
gtggtggaca ccgacctcat cgtcgacgac ggtggtgang nnacnctnct catccacnaa 120
ggcgtcaang ccnaggagcn cnatgagaag accggcgaa tcnccgannc caactccacc 180
ganaacgccg agctgcagat cgnngcttacc atcancagag anggggtgaa gaccganccc 240
atnaggnanc gcaagatgaa ggagcgtctc gttggggtnct ctgagggnac cancaactgga 300
gttaagagggc tcnatcag 318

<210> 3163
<211> 319
<212> nucleic acid
<213> Glycine max
<400> 3163

gtcgcangca cgcgtacgta agctcggaat tcggctcgag acggctgcga gaagacgaca 60
gaaggctcag cttgagttgt ggaaggagaa gagtaccggc aagtacgaga agaaagttta 120
cgttttgccc aagcaccttg atgagaaggt ggctgcactt caccttggca aacttggagc 180
taagctcacc aagcttagcc cggcccaggc tgattacatc agtgtgcctg ttgaggggtcc 240
ataaagcctg ctattacag gtactaagta attgagatta tcaacggaaa gtgagggaaa 300
gacaaaatcg gttttatga 319

<210> 3164
<211> 294
<212> nucleic acid
<213> Glycine max

<400> 3164

tcgcangcac gcntacgtaa gctcgggaatt cggctcgagc ggaaagtgag ggaaagacaa 60
aatcgggtttt atgaatcgga ttgattgttt aattttcctt tgataatctc aattacttag 120
tacctgtaat gagcaggctt gtatggaccc tcaacaggca cactgatgta atcagcctgg 180
gccgggctaa gcttgggtgag cttagctcca agtttgccaa ggtgaagtgc agccaccttc 240
tcatcaaggt gcttgggcaa aacgtaaact ttcttctcgt acttgccggt actc 294

<210> 3165

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 3165

nncgcatgca cgcgtacgta agctcgggaa ttcgggctcg agctctttct ctagtcctgt 60
tattttctcag cgcgtaaagc atgggctttg ttgggtggaga aaaccacgag tggtcgcgat 120
acaagggtcaa ggacctttcc caggccgact tcggccgctt cgagatcgag ctggccgagg 180
ttgagatgcc cggcctcatg gcctgtncgg accgagttcg gccctccca gccottcaag 240
ggggcccgca tcaccggctn cccttcacaca tgaccatcca gaccgccgtt ctca 294

<210> 3166

<211> 204

<212> nucleic acid

<213> Glycine max

<400> 3166

cgtcgcangc acgcgtacgt aagctcggaa ttcggtctga ggtttctgag gaaaccacca 60
ctggagttaa gaggctctat cagatgcagg cgaatgggac tcttctcttc cctgctatta 120
atgtcaatga ctctgtcacc aagagcaagt ttgacaactt gtatgggtgc cgtcactctc 180
tccctgatgg tctcatgagg gcta 204

<210> 3167

<211> 203

<212> nucleic acid

<213> Glycine max

<400> 3167

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gtcaccaaga gcaagtttga 60
 caacttgat ggggtgcgctc actctctccc tgatgggtctc atgaggggcta ccgatgttat 120
 gattgctgga aaggtggctg ttgtggctgg atatgggtgat gttggcaagg gttgtgctgc 180
 tgcaatgaag caggttggtg ctc 203

<210> 3168
 <211> 266
 <212> nucleic acid
 <213> Glycine max
 <400> 3168

gcaagtacga gaagaaagtt tacgttttgc ccaagcacct tgatgagaag gtggctgcac 60
 ttcaccttgg caaacttggg gctaagctca ccaagcttag cccggcccag gctgattaca 120
 tcagtgtgcc tgttgagggg ccatacaagc ctgctcatta caggtactaa gtaattgaga 180
 ttatcaacgg aaagtgaggg aaagacaaaa tcggnnttat gaatcggatt gattgtttta 240
 ttttcctttt tttgaatttt tgttgt 266

<210> 3169
 <211> 326
 <212> nucleic acid
 <213> Glycine max
 <400> 3169

ntctnntgna ngcgtacgta agctcggaat tnnctcgag ntcgagcngc gccgtcttcn 60
 anntggacag ggtgagaccc tccaggant actgggtggg caccgngcgc gccctcgact 120
 ggggccccgg tgggtgaccc gacctcatcg tnnacgacgg tggtgacgct acccttctca 180
 tccacgaagg cgtcaaggcc gaggagctct ntgagaagac cggcgaattc ccgancccaa 240
 ntccaccgac aagccggant ttcagatcgt gnttancatc atcagagatg gttgaagacc 300
 gttccaacca ggttacngca gatgaa 326

<210> 3170
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 3170

natcgatgca cgcgtacgta agctcggnat tcggctcgan ctcgagccga atcggtcga 60
 gggttgacca catgaagatn atganganca atgcnattgt anncaacatt ggnncacttt 120
 natcatnagn tcgacatnct nggggtggag nactaccccg gcgtgangcg catccaccat 180
 caagcccca accgacagat gggtcancce cgagaccaat gtcggcatca ttgtcttggc 240
 cgagggctcgt ttgatgaact tgggatgcgc cacaggacac cctagttttg tgatgtctgt 300
 cctcacnaac caggt 315

<210> 3171
 <211> 274
 <212> nucleic acid
 <213> Glycine max

<400> 3171

ncgcgtgnac gcgtacgtaa gctcggaatt cggctcgagg tttcctcacc actccctcca 60
 ttctctttct ctagtcctgt tatttctcag cgcgtaaagc atggctttgt tgtcnggaga 120
 anaccacgag tggtcgagag tacaaggtca aggaccttc ccaggccgac ttcgcccgcc 180
 tcgagatcga gctggccgag gttgngatgc ccggcctcat ggcntgtcgg accagagttcg 240
 gccctccca gcccttcaag ggggcccgca tcac 274

<210> 3172
 <211> 282
 <212> nucleic acid
 <213> Glycine max

<400> 3172

gtgcgancga cgcgtncgga ngcacgctcn ctttgcctta gtgcctgtta tttctcanen 60
 cgtaaagcat ggctttgttg gtggagaaaa ccacnagtgg tgcgcgagta canggtinnag 120
 gacctttgcc caggccgact tcnccgcct cgagatcgag ctggccgagg ttganatgcc 180
 cggcctcatg gcctgttcgg accgagttcg gccctccca ncccttcaag ggggcccgca 240
 tcaccggctc cctccacatg accatccaga ncgcccgttct ca 282

<210> 3173
 <211> 312
 <212> nucleic acid
 <213> Glycine max

<400> 3173

acgtcgcang cagcntacg taagctcgga attcggctcg agtacgtttt gccaagcac 60
 cttgntgaga aggtggctgc acttcacctg ggcaaacttg gngctaagct gaccagctt 120
 agcaagtccc aggttgatta catcagtgtg cctgttgagg gtccatacaa gcctgtcac 180
 tacaggctact aagtgattga gatgatcaac tgaaaagtga gtgagggaaa gacaaaaatc 240
 ggttttatca atcggatttg attgtttaat tttccttttt tgatttttgg tgtagactt 300
 tcagatttgn gg 312

<210> 3174

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3174

angnacangc acgcgtacgt aagctcgga ttcggctcga gtgccaagc accttgatga 60
 gaaggctggct gcaattcacc tgggcaaact tggngctaag ctgaccacgc ttagcaagtc 120
 ncaggctgat tacatcagtg tgctgttga ggtccatac aagcctgtc actacaggta 180
 ctaagtgatt gagatgatca actgaaaagt gaggtaggga aagacaaaaa tcggttttat 240
 caatcggatt tgattgttta attttccttt ttttgatttt tgggtgttaga cttttca 297

<210> 3175

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3175

tcgcntgcac gcgtacgtaa gctcgggaatt cggctcgagt ttacgttttg cccaagcacc 60
 ttgatgagaa ggtggctgca cttcanctgg gcaaacttgg acctaagctg acccagctta 120
 gcaagtccca ggctgattac atcagtgtgc ctgttgaggg tccatacaag cctgtctact 180
 acaggctacta agtgattgan atgntcaact gaaaagtgag tgaggggaaag acaaaaaatcg 240
 nttttntcaa tcggatttga ttgtttaatt ttcctttttt tgatttttgg tgtaga 297

<210> 3176

<211> 289

<212> nucleic acid

1118
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<213> Glycine max
 <400> 3176
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 aagggtggctg cacttcacct gggcncactt ggngctaagc tgacccagct tgcncaaatc 120
 ccaggctgat tacatcagtg tgccctgttg gggctccatac aagcctgctc actacaggta 180
 ctaagtgatt gagatgatca actgaaaagt gaggtaggga aaggacaaaa atcgggtttta 240
 tcaatcggat ttgatgttta attttccttt tttgattttg gtgtttagan 289

<210> 3177
 <211> 336
 <212> nucleic acid
 <213> Glycine max
 <400> 3177
 gngangcagn gtacgtaagc tcggaattcg gctcgaggag agagagagag agagagagag 60
 atctatctat ctatcaagat ngcggtgttg gttgaaaaaa aaaannattg anaggganta 120
 caagggtgaag ganatgatgc aagccgnttt nggaagattg gaaattcgag ctggcgagg 180
 ttgaaatgcc cggcctcatg tccncccg accgagttcg gcccctcttc aatccttcaa 240
 gggcgctagg atcancggct cctccacat gaccatcnan agccgncgtc cttcatngag 300
 acnctcaccg ctctcggcgc cgaggtccgc tgggtgc 336

<210> 3178
 <211> 209
 <212> nucleic acid
 <213> Glycine max
 <400> 3178
 tnattacnag tacgnaagct cggaattcgg ntcgagccct ccaggagtag tgggtggtgca 60
 ccgagcgcg cctcgactgg ggcncggtg gtggaccga ccttcacgt cgacgacgg 120
 ggtgacgnta cccttctcat ccacgaaggc gtcaaggncg agganctcta tgagaanacc 180
 ggcgaaactg ccganccan ctccacaaa 209

<210> 3179
 <211> 291
 <212> nucleic acid

<213> Glycine max

<400> 3179

nagtcgcang cacgcgtacg taagctcgga attcggctcg aggctcacca actcccgtc 60
ccatttcctt atttatagac agagtctgat tgtttcctca ccactccctc cantctcttt 120
ctcctagtcc tgttatttct cagcgcgtaa agcatggctt tgttggtgga gaaaaccacg 180
agtggctcgc agtacaaggt caaggacctt tcccaggcgc attcgggcgc cctcgagatc 240
gagctggcgc aggttgagat gcccggcctc atggcctgtc ggaccgagtt c 291

<210> 3180

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3180

nacgtgcgat gcacgcgtac gtnagctcgg aattcggctc gagtgnngaag accggggaac 60
tccccgaccc taactccact gacaacgcgc agttccagat cgtgcttacc atcatacaga 120
gatgggttga agaccgatcc caccaggtac cgcaagatga aggagcgtct cgttgggggtt 180
tctgaggaaa ccaccactgg tgtaagagg ctatatcaga tgcaggcgat tgggntntat 240
ttccccgctna taataatnnc nngnnntctg ttaccaagng cngtntnaca acttgnc 297

<210> 3181

<211> 208

<212> nucleic acid

<213> Glycine max

<400> 3181

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tacgttttgc ccaagcacct 60
tgatgagaag gtggctgcac ttcacctggg caaacttggg gctaagctga cccagcttag 120
caagtcccag gctgattann ncagtgtgcc tgttgagggt ccatacaagc cgctcactac 180
agggtactaag tgattgagat gatcaact 208

<210> 3182

<211> 212

<212> nucleic acid

<213> Glycine max

<400> 3182

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctctttct ctagtcctgt 60
tattttctcag cgcgtaaagc atggccttctg tggaggagaa aaccacnagt ggtcgcgagt 120
acaagggtcaa ggacctttcc caggccgact tcggcngcct cgagatcnag ctggccgagg 180
ttgagatgcc cggcctcatg gctgtcgga cc 212

<210> 3183

<211> 317

<212> nucleic acid

<213> Glycine max

<400> 3183

aagtncncat gcaagctac gtaantcgga attcggctcg agctctagtc ctgttatttc 60
tcancgcgta aagcatgggc ttgtttggtg gagaaaacca cgagtngtcc gctagtacaa 120
ggtcaaggac ctttcccagg ccgacttcng ccgcctcgag atcgagctgg ccgagggtga 180
natgcccggc ctcatggcct gtnggaccga ntccggcccc ttcccaaccc ttcaaggggg 240
cccgatcan cggctccctn canatganca tccagaancg cgttntcatt gngaccctna 300
ncggctttgg ggcggag 317

<210> 3184

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 3184

tcgcangcag ncgtacgtaa gctcggaatt cggctcgang anaggtgnct gcacttcacc 60
tgggcaaact tggcnctaag ctgaccanc ttagcaagtc ccaggctgat tacatcagtg 120
tgcncgttga gggccatac aagcctgctc antannngta ctaagtgatt gagatgatca 180
actgaaaagt gagtgaggga aagacaaaaa tcggttttat caatcggatt tgattgttta 240
attttccttt ttgattttt ggtgttngac ttttcagaat gtggtagaag aatt 294

<210> 3185

<211> 245

<212> nucleic acid

<213> Glycine max

<400> 3185

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tgtccctccc ggcaccngt tcggccctc ttcaaccctt caagggcgcn angatnaccg 180

gntccctcca caatgagcnn ncaaanagcc gtacctnaaa cgnagacncg cacnngccng 240

ggggc 245

<210> 3186

<211> 234

<212> nucleic acid

<213> Glycine max

<400> 3186

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atcggtcga ctttctctag tctgtttatt tctcagcgcg taaagcatgn ctttgttggt 120

ggagaaaaca nacgagtggg cgcgagtaca aggtcaagga cctttcccag gccgacttcg 180

gccgntcga gatcgagctg gccgaggttg agatgcccg cctcatggcc tgtn 234

<210> 3187

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 3187

tcgcnnngcac gcgtacgtna gctcgnantt cggcnccgag tggnaggagg taaggctggg 60

tcgaccaga tctagttgag ctcaccaact cccgctccca tttccttatt tatagacaga 120

gtctgattgt ttctcacca ctccctccan tctctttctc tagtcctgtt atttctcagc 180

gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 240

gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgc 298

<210> 3188

<211> 221

<212> nucleic acid

<213> Glycine max

<400> 3188

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 cgctcgagcc ggtcctgtna tntctcagcg cgtaaagcat ggctttantt ggtgganaaa 120
 accacgagtg gtcgcgagta caaggtcaag gacctttccc aggccgactt cgggccccctn 180
 cgagatcgag ctggccgagg ttgagatgcc cggcctcatg g 221

<210> 3189
 <211> 291
 <212> nucleic acid
 <213> Glycine max
 <400> 3189

anncananaa tnatgcacgc gtacgtaagc tcggaattcg gctcgagatt gtttcctcac 60
 cactccctcc antctctttc tctagtcctg ttattttctca gcgcgtaaag catggctttg 120
 ttggtggaga aaaccacgag tggtcganag taanaaggctc aaggactttc ccaggccgac 180
 ttcgnggcc tcgagatoga gctggccgag gttnaaatgc cgggcctcat ggctggncgg 240
 acgattnggg cccctcnaa cctttaaggg gggccnaaat cangggntcc n 291

<210> 3190
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 3190

ncgcgatgca gcgtacgtaa gctcggaatt cggctcgagg ttattttctca gcgcgtaaag 60
 catggctttg tnggtggaga aaaccacgag tggtcgcgag tacaagggtca aggacctttc 120
 ccaggccgac ttcgcccgcc tcgagattcg agctggncgg aggttgagat gcccgacct 180
 catggcctgt ncggaccgag ttnggncccc taccagccc tttcaagggg gncccgcatc 240
 accggcnccc nccacatgna ccatccagtg ccgnccgttg ttcattgn gn accctgcacc 300
 gcc 303

<210> 3191
 <211> 144
 <212> nucleic acid
 <213> Glycine max
 <400> 3191

ngcaggcacg cgtacgtaag ctncggaatt cggctcgagn cggctcgagg ggttggtgctg 60
 ctgcattgaa gcaggctggg gctcgtgtca tcgtgactga gattgacccc atttggtgcc 120
 ttcaggctct catggaaggg cctt 144

<210> 3192
 <211> 134
 <212> nucleic acid
 <213> Glycine max
 <400> 3192

aacgtcgcat gcacgcgtac gtaagctcgg aattcggctc gagcccgacc tnatcgtcga 60
 cgacggtggg gacgtaccc ttctcatgcc acgaaggcgt tnaggccgag gagctctatg 120
 agaagaccgg cgaa 134

3193
 303
 nucleic acid
 Glycine max
 3193

<210> 3193
 <211> 303
 <212> nucleic acid
 <213> Glycine max
 <400> 3193

acgtcgcatg cagcgcgtacg taagctcggg attcggctcg aggccttagca agtcccaggg 60
 ctgattacat cagtgtgcct gttgaggggc catacaagcc tgctcactac aggtactaag 120
 tgattganat gatcaactga aaagtgagtg agggaaagac aaaaatcggg tttatcaatc 180
 ggatttgatt gtttaatttt cctttttttg atttttgggt ttagactttt cagatttgtg 240
 gtagaagaat gtagccattt ttatttctgt agaacttttg ttcgggtgggt gggaccagta 300
 agg 303

<210> 3194
 <211> 315
 <212> nucleic acid
 <213> Glycine max
 <400> 3194

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 gtgtgccacg ggacacccca gctttgtgat gtcgtgctcc ttcaccaacc aggtcatngc 120
 tcagcttgaa ttgtggaaag agaagggttc tgggaagtat gagaagaagg tgnatgtgtt 180

gccaagcac cttgacngga aagtggcttc tctccacctt ggccagnttg gagctaggct 240
 caccaagctt tccanagacc aagctgatta catcagtgtg cctgttgagg gtccatacaa 300
 gccgctccnc acagt 315

<210> 3195
 <211> 290
 <212> nucleic acid
 <213> Glycine max

<400> 3195

cancacactc gcangcacgc gtacgtaagc tcggaattcg gctcgagcag gtcattgctc 60
 agcttgaatt gtggaaagag aagggttctg ggaagtatga gaagaagggtg tatgtgttgc 120
 ccaagcacct tgacgagaaa gtggcttctc tccaccttgg ccagcttggg gctaggctca 180
 ccaagctttc caaagaccaa gctgattaca tcagtgtgcc tgttgagggt ccatacaagc 240
 ctgctcacta caggtactga tccatcctat tgggggagaa taaacctaaa 290

<210> 3196
 <211> 217
 <212> nucleic acid
 <213> Glycine max

<400> 3196

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 gagaaggggt ctgggaagta tgagaagaag gtgtatgtgt tgccaagca ccttgacgag 120
 aaagtggctt ctctccacct tggccagctt ggagctaggc tcaccaagct ttccaaagac 180
 caagctgatt acatcagtgt gcctgttgag ggtccat 217

<210> 3197
 <211> 255
 <212> nucleic acid
 <213> Glycine max

<400> 3197

gaaagagaag ggttctggga agtatgagaa gaagggtgat gtgttgccca agcaccttga 60
 cgagaaagtg gcttctctcc accttggcca gcttggagct aggctcacca agctttccaa 120
 agaccagctg attacatcag tgtgctgttg angggggcca taanagcttg tcnctnangg 180

nnnnggnccn ncctttgggg gggaannaac ccgaantntn ttnattcgg ggggggnttg 240
 tnnantttnn ttng 255

<210> 3198
 <211> 338
 <212> nucleic acid
 <213> Glycine max
 <400> 3198

aggaccatgc cgccgccgcc atcgcccgcg acaggcctcc gtcttcgcct ggaagggatga 60
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 cccgatctca tcgtcgacga cggcggcgac accactcttc tcattcacga gggcgtcaag 180
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 gattncgat cgtgctgagc atcatcaggg atggttgaag accgatccca agaggtacca 300
 caagatgaag acagaatcgt cgggtgtctcc gaagaaac 338

<210> 3199
 <211> 317
 <212> nucleic acid
 <213> Glycine max
 <400> 3199

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 anctcatcgt ccgacgacgg cggcgacacc actcttctca ttcacgaggg cgtcaaggcc 180
 gaggagatct ttgagaagac cggccagttc cccgaccggg ctccctccga caatgcggag 240
 ttccagatcg tgctgagcat cattcagggg tggcttgaag accgatccca agaggtacca 300
 caagatgaag gacagaa 317

<210> 3200
 <211> 290
 <212> nucleic acid
 <213> Glycine max
 <400> 3200

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ggaggtgacg gatcttgaca tggttgatgc tncatagta gaagggaaaa caaaagtggc 120
 ttacttcga atctgtttcc aacccacccc ttacgggtgc gaacatacct gaactgtgcc 180
 acatggcaca ccggaagga gtgacggtgg tggaggacaa cacgttcgcg cccatggtgc 240
 ttccgccagc gcgtcttggg gctgatgttg tggttcacag tatctccaag 290

<210> 3201
 <211> 213
 <212> nucleic acid
 <213> Glycine max
 <400> 3201

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 atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaacccca 120
 cccttacggt tgcgaacata cctgaactgt gccacatggc acaccggaag ggagtgcg 180
 tggagggtga caacacgttc gcgcccatgg tgc 213

<210> 3202
 <211> 297
 <212> nucleic acid
 <213> Glycine max
 <400> 3202

cncangcacg cgtacgtaag ctccgaattc ggctcgaggt gggacccacg cgtccctctc 60
 acactttctc ccgcgcacgt gcggaatatc cactagcttc gtggagggtga cggatcttga 120
 catggttgat gctgccatag tagaagggaa aacaaaagtg ctttacttcg aatctgtttc 180
 caacccacc cttacggttg cgaacatacc tgaactgtgc cacatggcac accggaaggg 240
 agtgacggtg gtggaggaca acacgttcgc gcccatggtg ctttcgccag cgcgtct 297

<210> 3203
 <211> 300
 <212> nucleic acid
 <213> Glycine max
 <400> 3203

gtcncnngta cgtaanctcg gaattcggct cgagcngaca ancccaannc ccaagcccaa 60
 caatctgcat ccccgccgc ggccgtgca accaaatggg ccgtggacag ctggaagtcc 120

aagaaggccc tgcagntgcc cgaatacccc aaccaggagg atctcgaggc cgtcctccgc 180
 accctcgacg cntnccccctc anategtctt cgcggcgag gcccgganac tcgaggagca 240
 cctcgccgag gccgccatng gaaatgcntt cttecnenan ggcgnagatg tnccnagagt 300

<210> 3204
 <211> 434
 <212> nucleic acid
 <213> Glycine max
 <400> 3204

1127

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 cacngcgaac anggtggtcg nctccagaac tctntacngt gggacccacn cgtcctctc 120
 acactttctc ccgcgcacgt gcggaatatc cactagcttc gtggaggtga cngatcttga 180
 catggnttat gctgccataa tagaaaggaa aacaaaagtg ctttacttnt aatctggttc 240
 caaccccacc cttacngttg cgaacatacc tgaactgtgc cacatggcac accggaaggg 300
 agtgactgtg gtggtggaca acacgttcgt gcccatggtg ctttcgccag cgcgtntttg 360
 gtgcttatgt ttgtncctca cagtatctcc aagttcatna atnggtgggg cccgatatta 420
 ttgcangagc ggng 434